

Virginia Arboviral Surveillance 2019

Virginia Mosquito Control Association

Virginia Beach

January 28th 2020

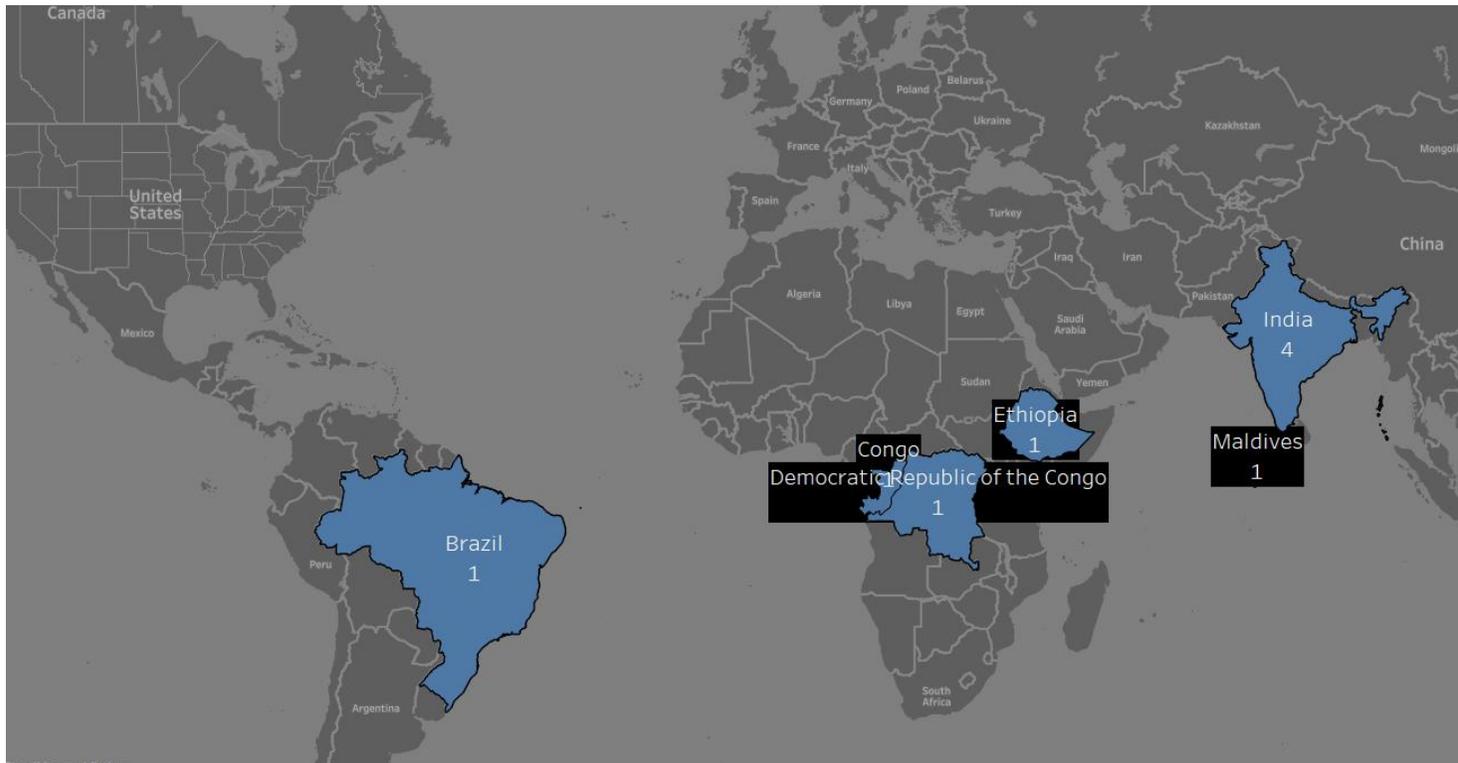
VA Human Infections from Arboviral Diseases

Arboviral Condition	2019	3 yr avg. trend	3 yr avg.	2018	2017	2016
La Crosse virus, neuroinvasive	0		1	2	0	0
West Nile infection, neuroinvasive	4		19	38	12	6
West Nile infection, non-neuroinvasive	2		5	10	1	2
Chikungunya virus diseases	9		5	3	5	6
Dengue virus	19		16	8	12	27
Zika virus, disease	1		39	1	6	109
Grand Total	62			73	44	172

Imported Arboviral Disease Cases in VA in 2019

Chikungunya (9 travel-related cases)

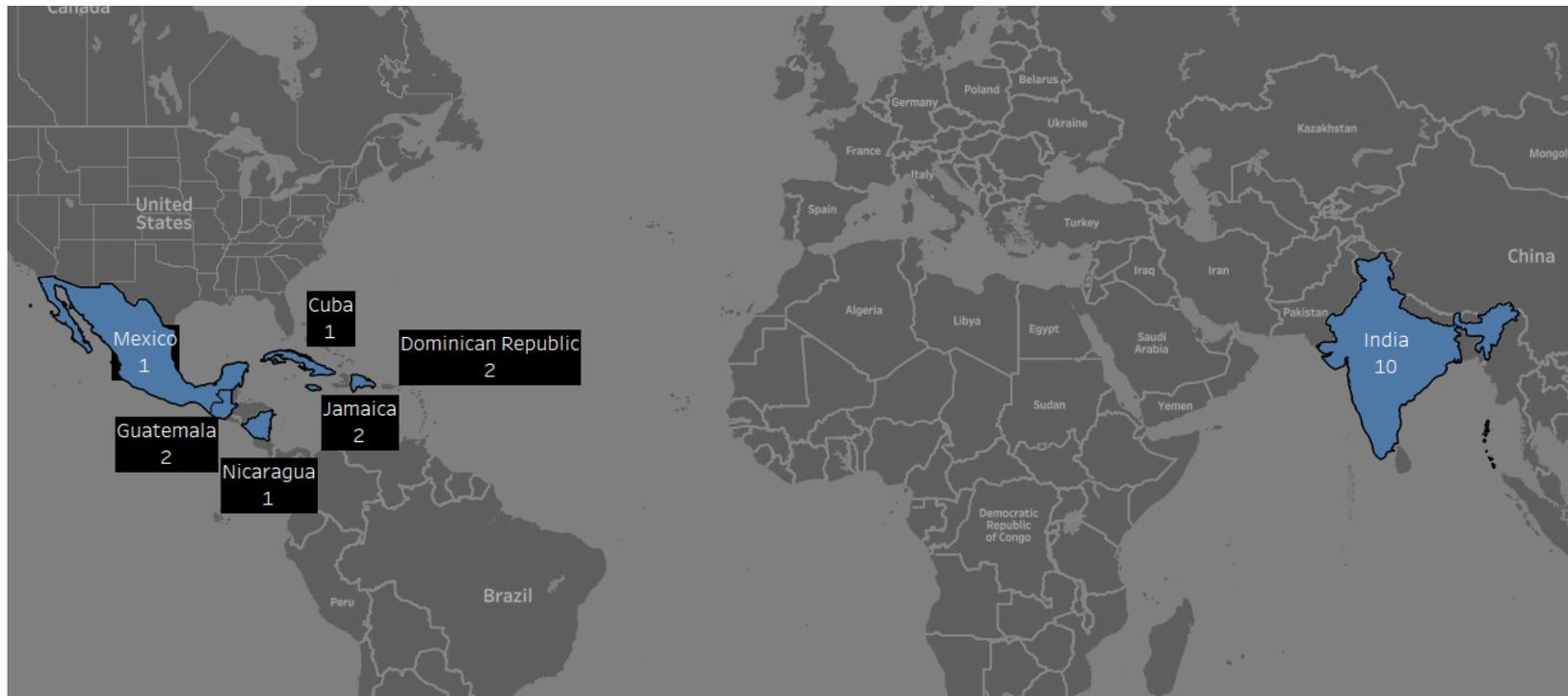
- India (4), Brazil (1), Congo (1), DRC (1), Ethiopia (1), Maldives (1)



Imported Arboviral Disease Cases in VA in 2019

Dengue (19 travel-related cases)

- India (10), Guatemala (2), DR (2), Jamaica (2), Cuba (1), Mexico (1), Nicaragua (1)



Imported Arboviral Disease Cases in VA in 2019

Zika (28 travel-related cases)

- Honduras (13), El Salvador (10), Guatemala (2), India (3)



West Nile Virus in US in 2018

As of Jan 8th, 2,544 WNV human cases found in 48 states and DC.

- 447 more WNV cases than 2017, 395 more than 2016
- 63% classified as neuroinvasive (meningitis or encephalitis)
- 37% classified as non-neuroinvasive (febrile illness)
- 357 WNV presumptive viremic donors reported from 35 states
- 137 fatalities (5.3% CFR) reported from 35 states

Figure 3. West Nile virus (WNV) neuroinvasive disease incidence* reported to ArboNET, by state — United States, 2018 (as of January 8, 2019)

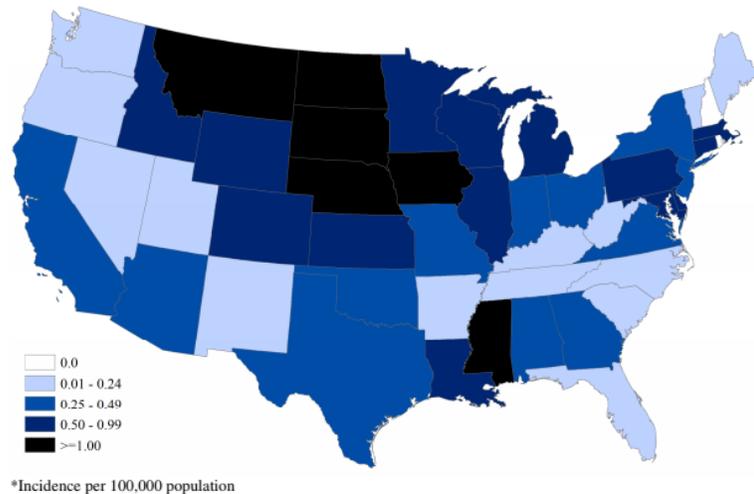
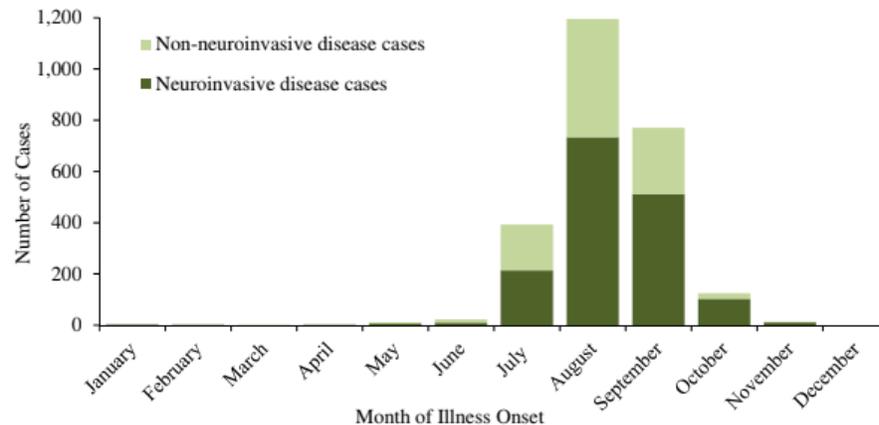


Figure 2. West Nile virus disease cases reported to ArboNET, by month of onset* — United States, 2018 (as of January 8, 2019)

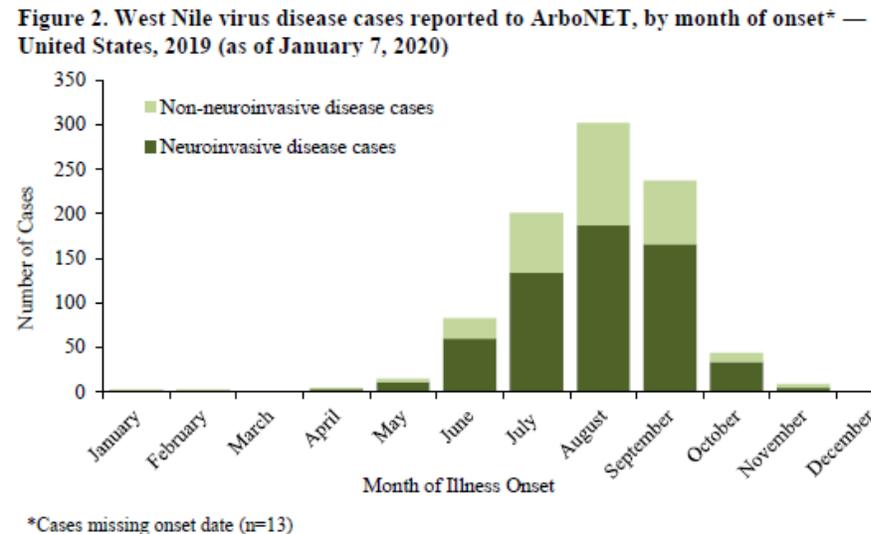
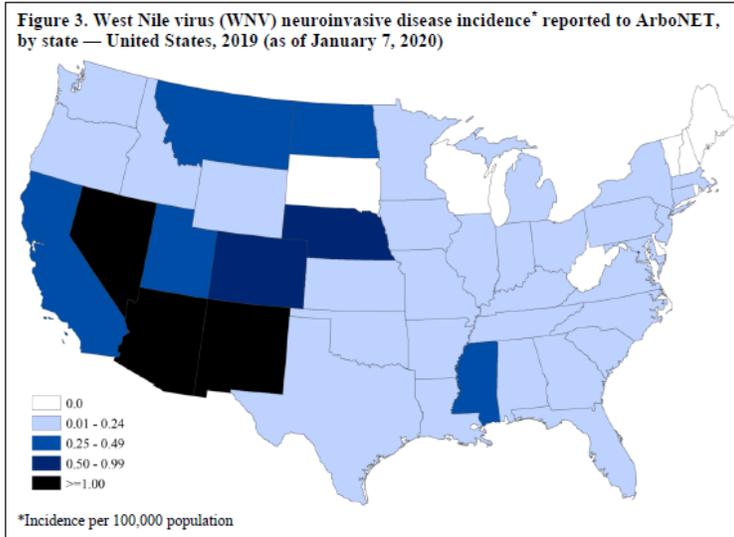


*Cases missing onset date (n=2)

West Nile Virus in US in 2019

As of Jan 7th, 917 WNV human cases found in 43 states and DC.

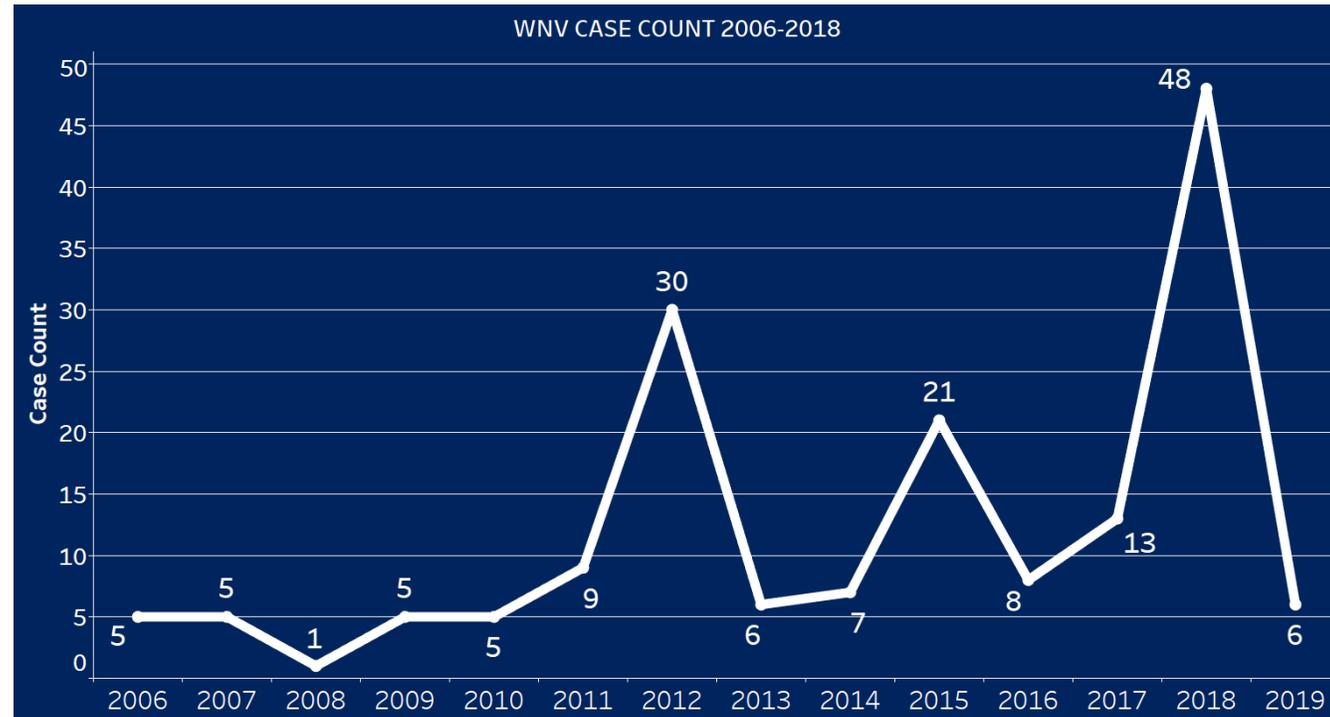
- Over 60% decrease from 2018 cases
- 66% classified as neuroinvasive (meningitis or encephalitis)
- 34% classified as non-neuroinvasive (febrile illness)
- 100 WNV presumptive viremic donors reported from 23 states
- 51 fatalities (5.5% CFR) reported from 35 states



West Nile in VA in 2019

6 WNV human cases in 4 jurisdictions

- 66% (4 cases) neuroinvasive
- 33% (2 cases) non-neuroinvasive
- 1 fatality reported

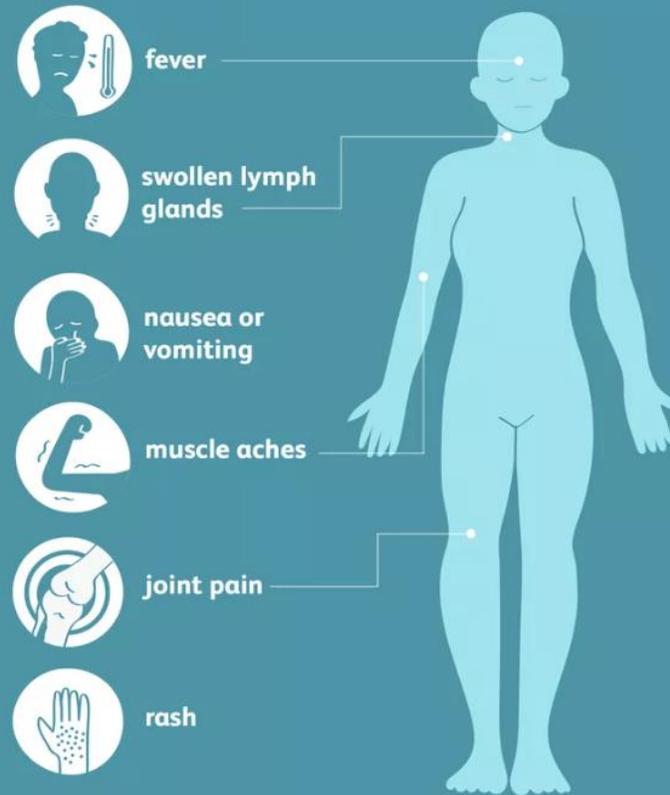


2018-2019 VA WNV CASES (n=54, 42 neuro & 12 non-neuro)

West Nile Virus

Common Symptoms

- 98% reported fever
- 65% reported headache
- 61% reported altered mental status
- 33% reported stiff neck
- 50% reported nausea/vomiting
- 48% reported muscle weakness
- 26% reported arthralgia (joint pain)
- 30% reported rash



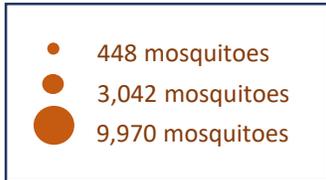
17% (n=9) reported as fatal

38/54 encephalitis/meningitis

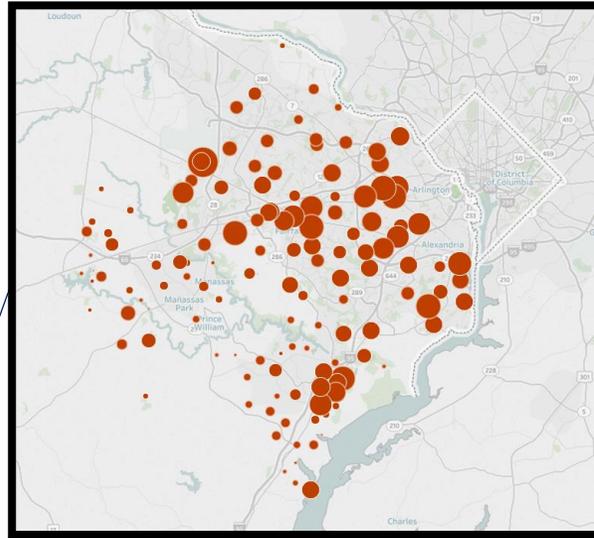
verywell

**54 cases total (42 neuro, 12 non-neuro)

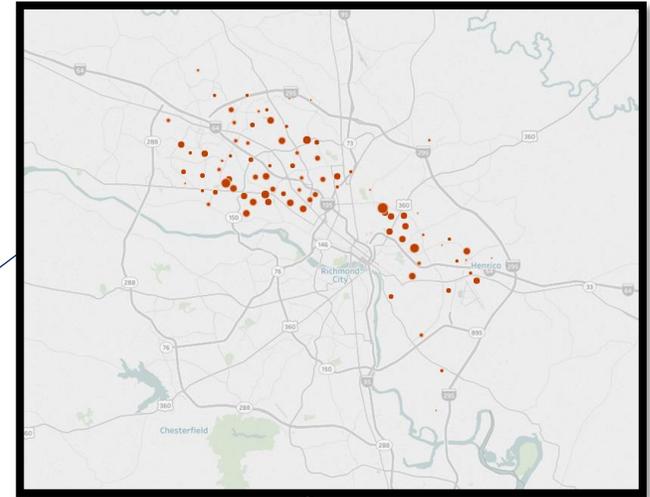
There were **316,021** mosquitoes trapped and tested in **9,166** pools during 2019 surveillance.



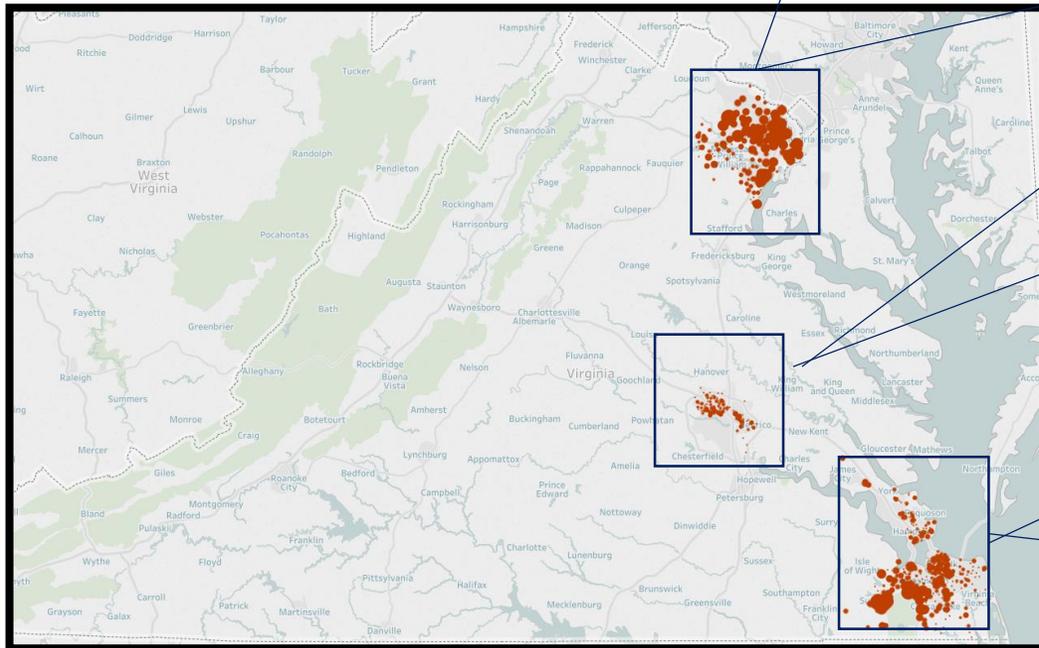
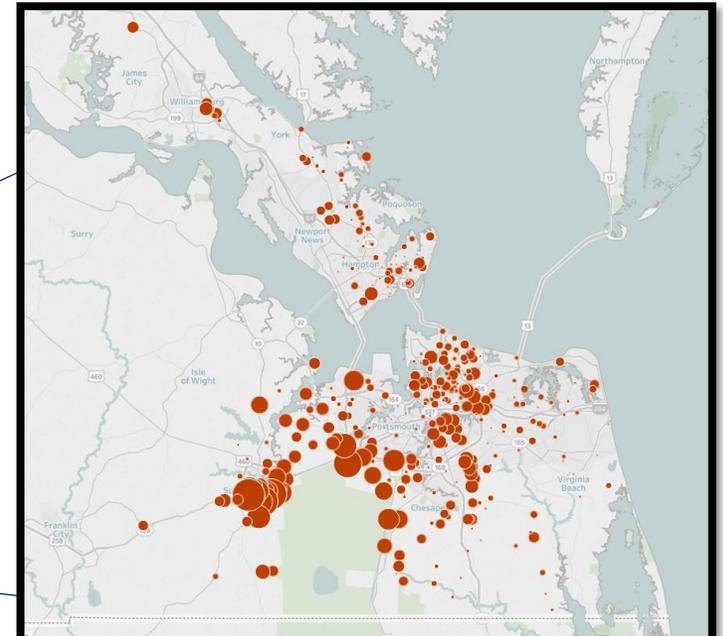
NOVA



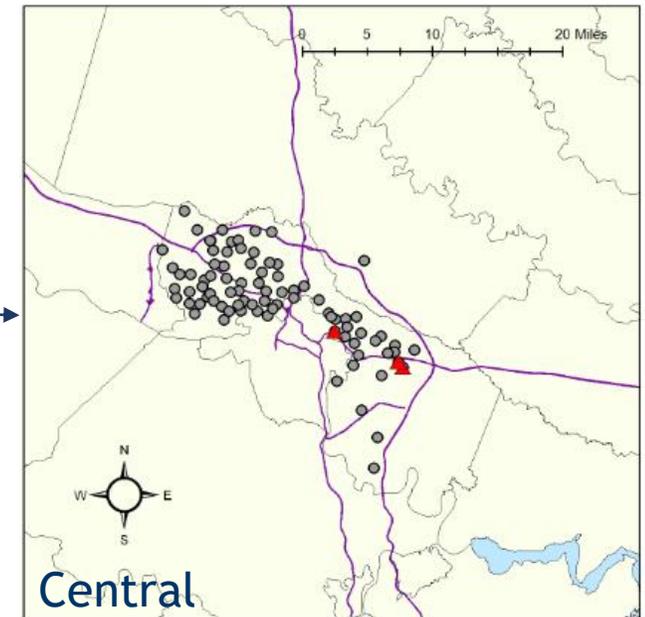
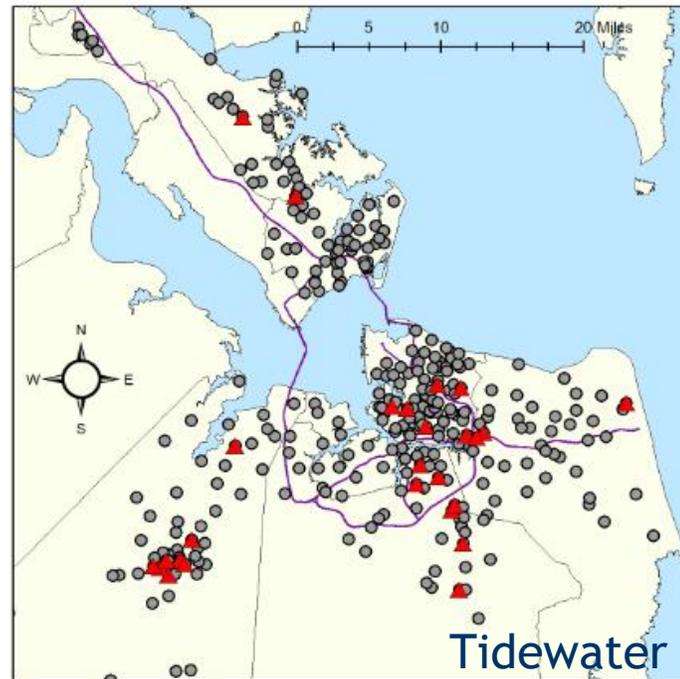
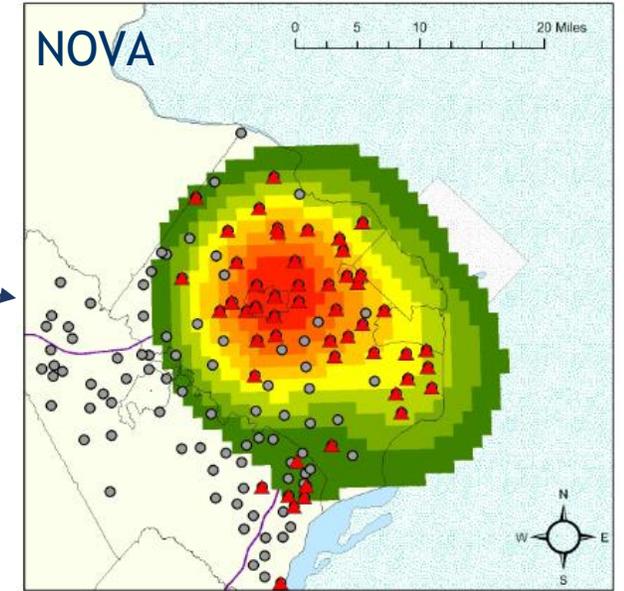
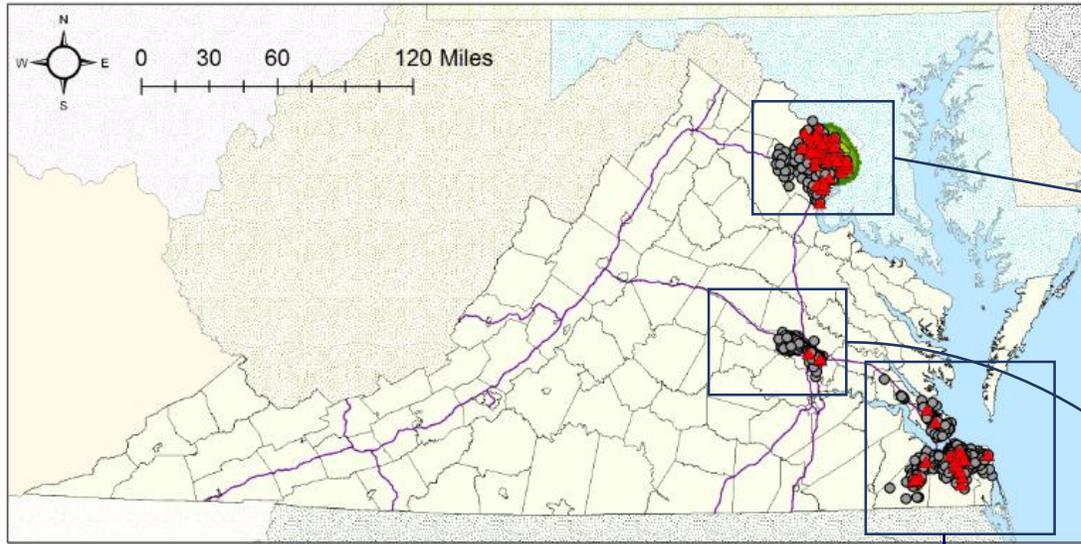
Central



Tidewater



Of 8,501 pools, 1.99% (n=169 pools) were WNV positive.

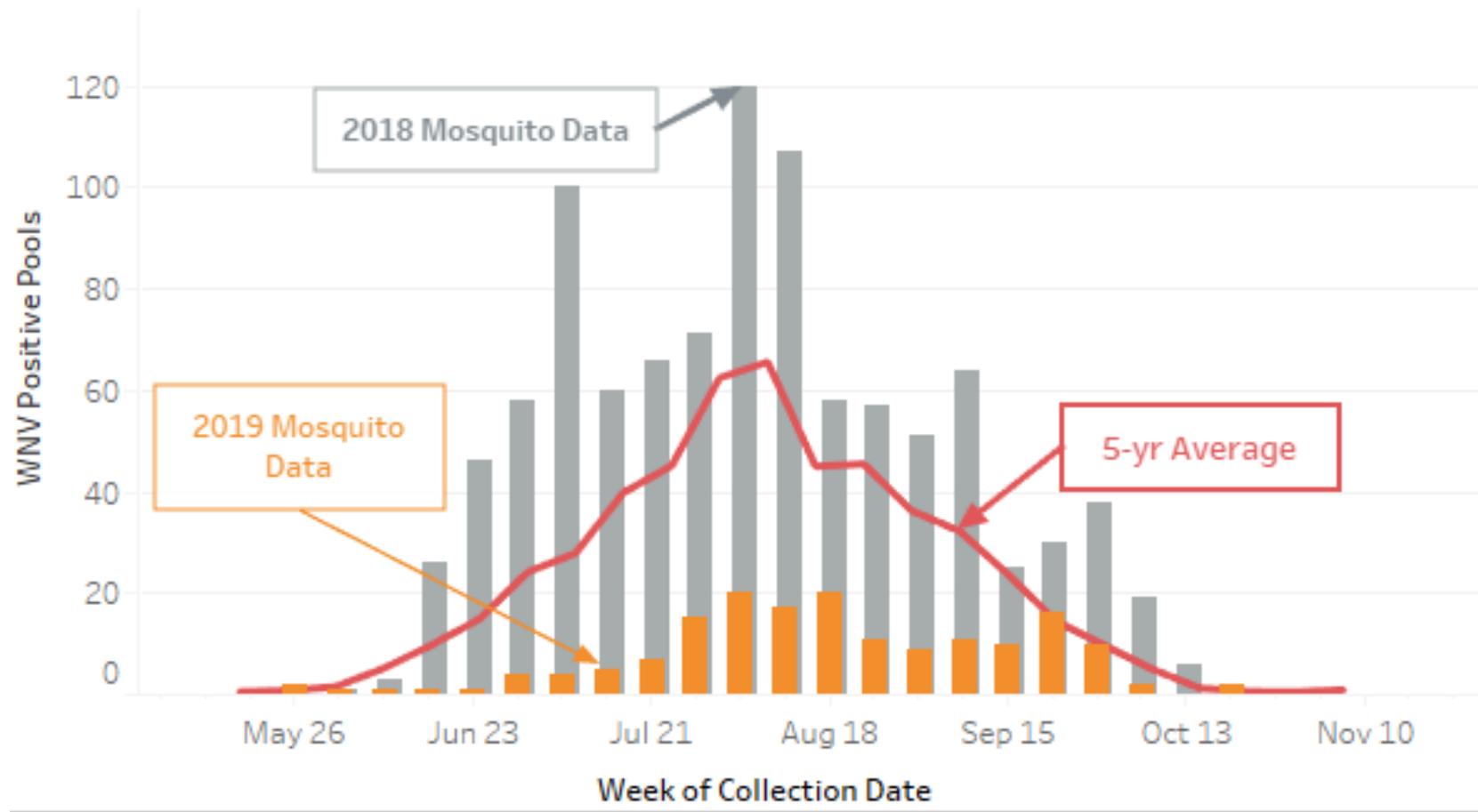


WNV Testing Mosquito Pool Sites

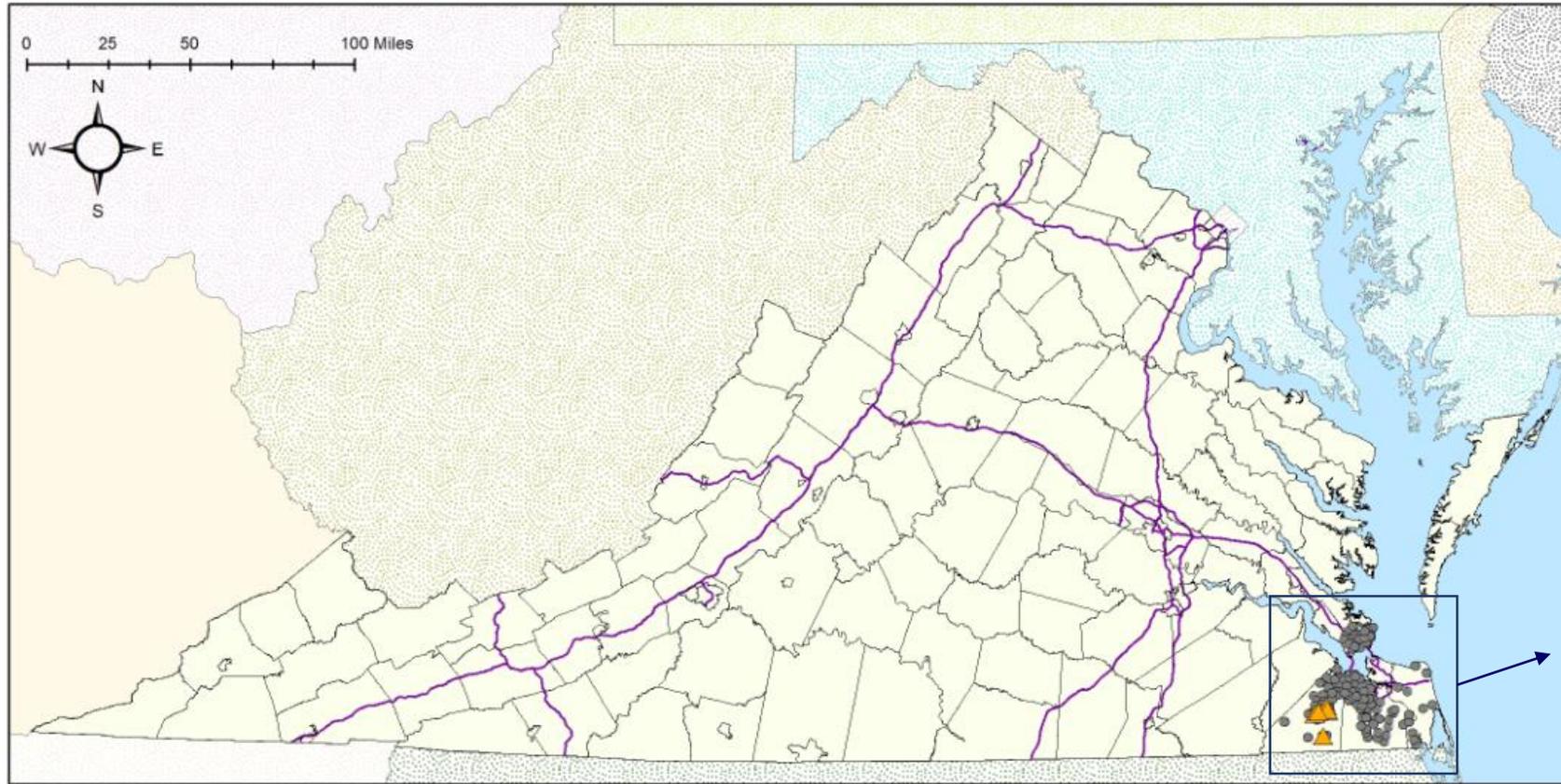
- ▲ WNV Pos
- WNV Neg

Of 8,501 pools, 1.99% (n=169 pools) were WNV positive.

West Nile Virus Positive Mosquito Pools

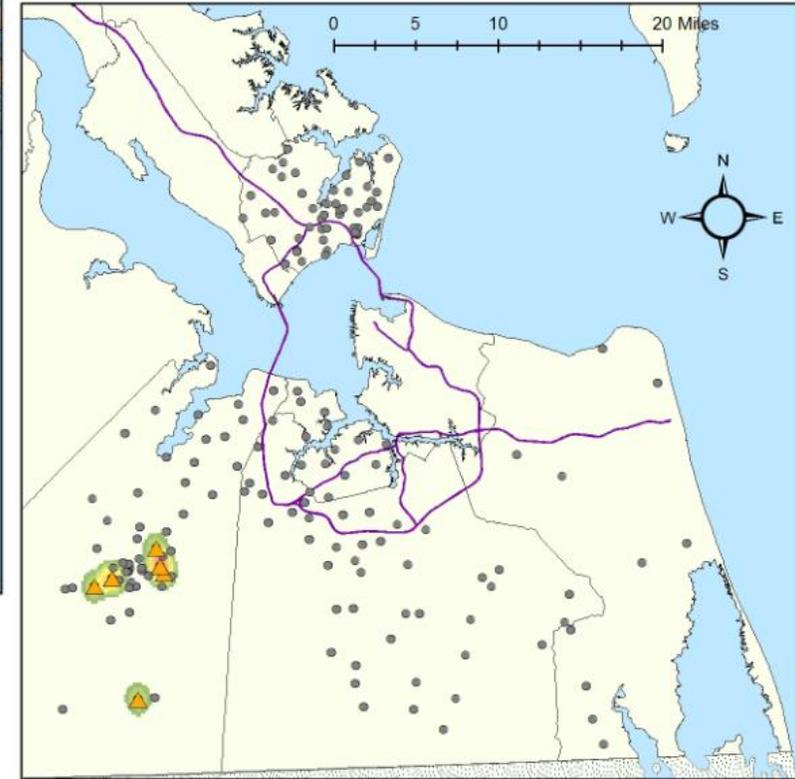


Of 2,051 pools, 0.39% (n=8 pools) were EEE positive.



EEE Mosquito Testing Pool Sites

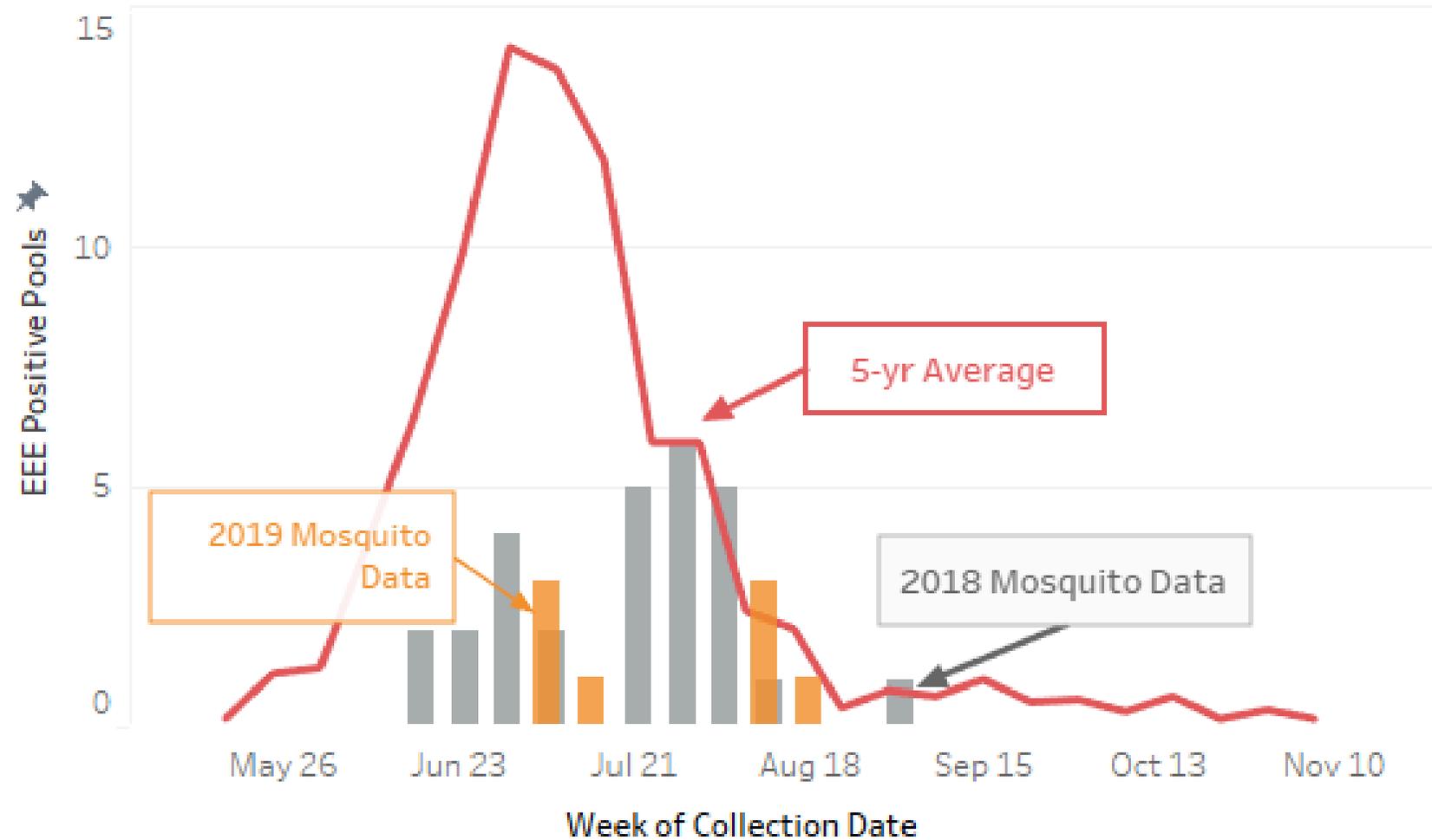
- ▲ EEEPos
- EEE

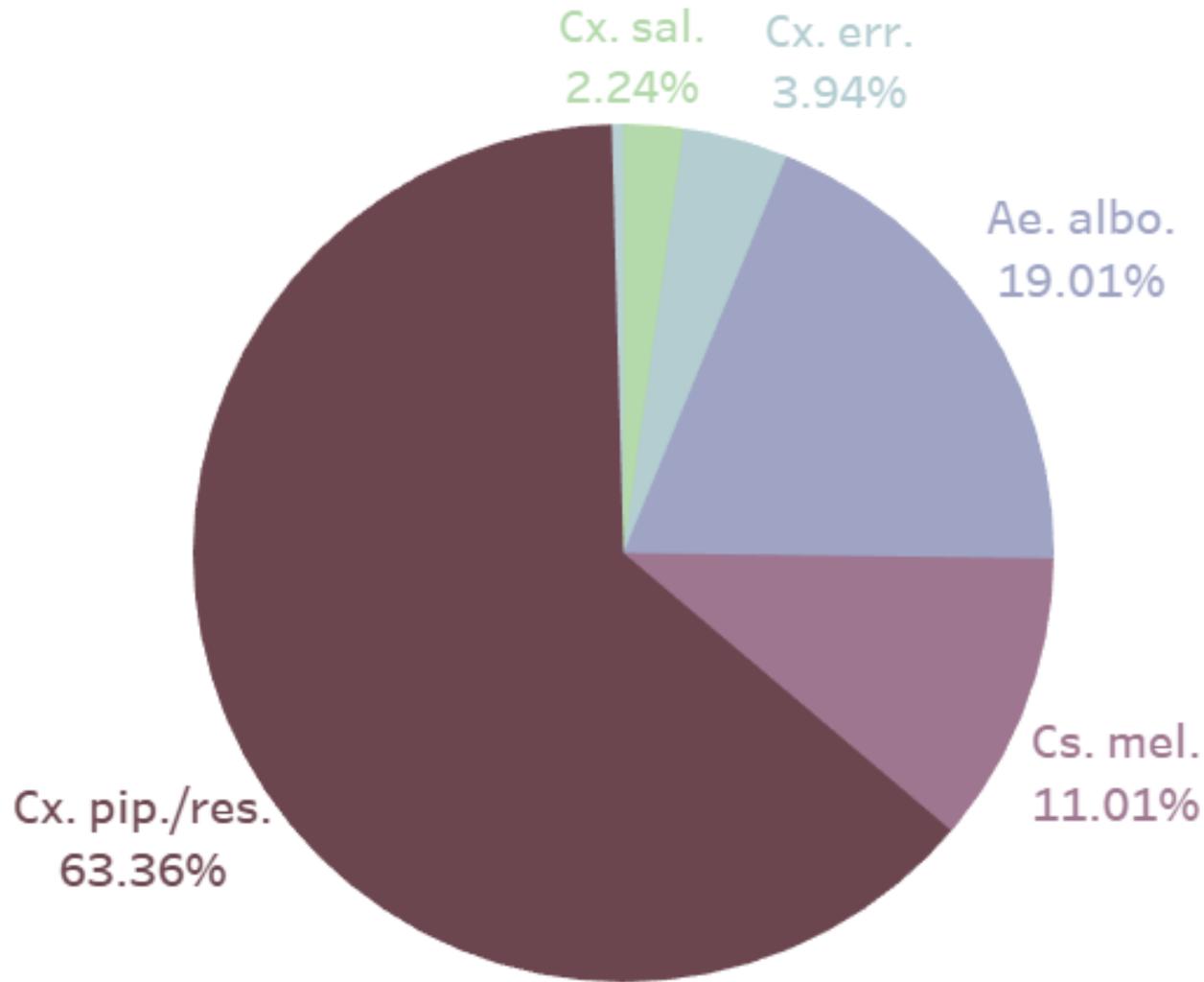


Tidewater

Of 2,051 pools, 0.39% (n=8 pools) were EEE positive.

Eastern Equine Encephalitis Positive Mosquito Pools



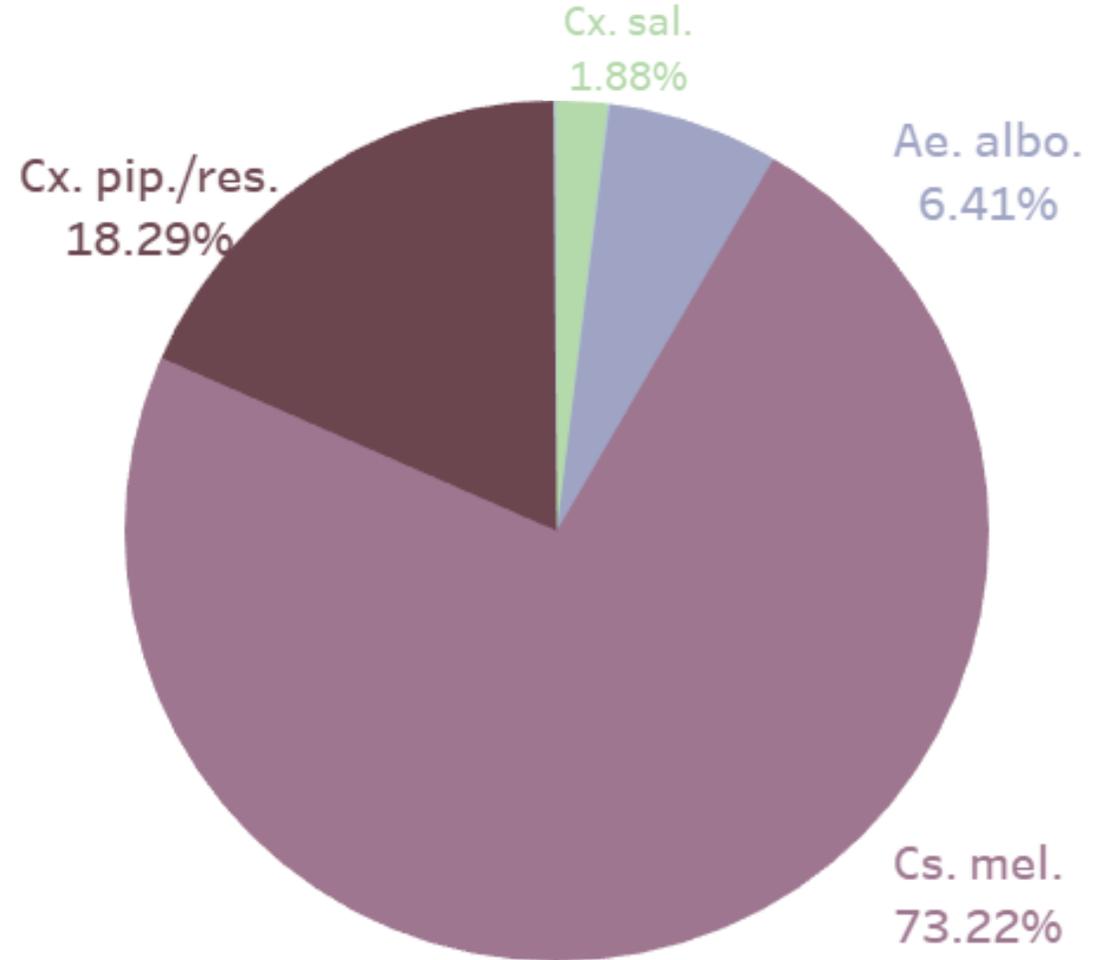


MOSQUITO SPECIES TESTED FOR WNV

(2019, n=8,501 pools)

165 *Cx. pip./res.* pools WNV+

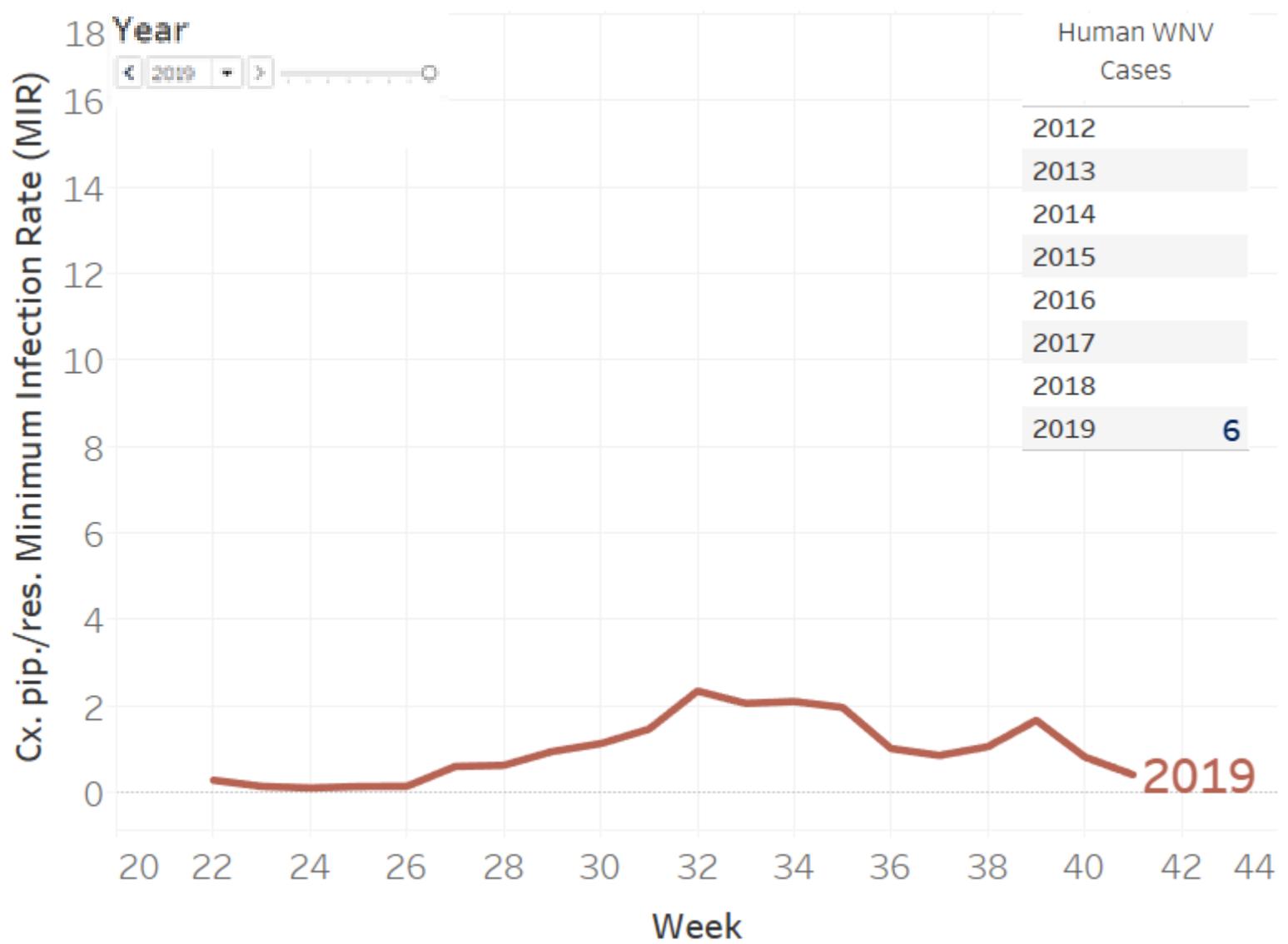
4 *Cs. mel.* pools WNV+



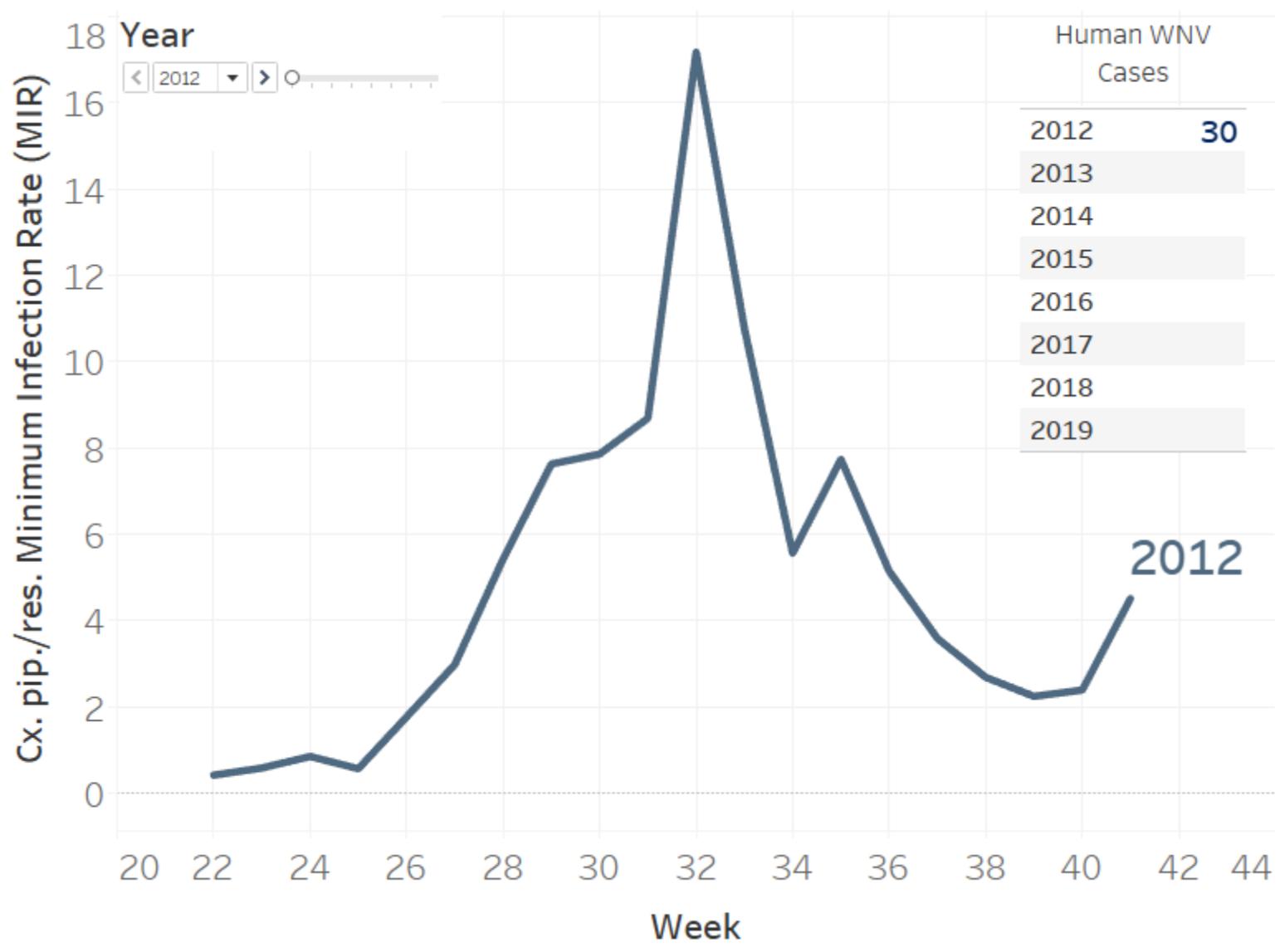
MOSQUITO SPECIES TESTED FOR EEE

(2019, n=2,051 pools)

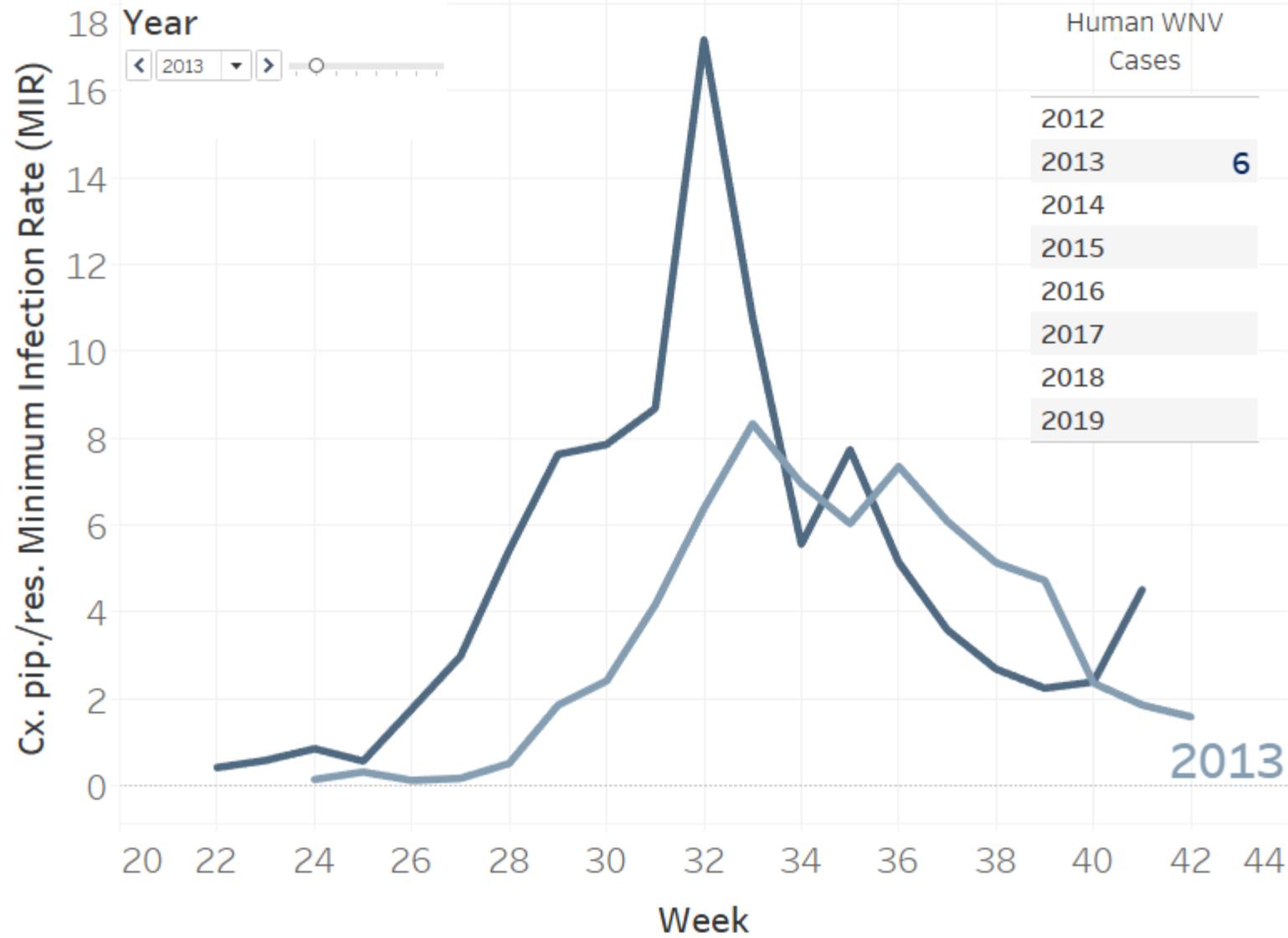
8 *Cs. mel.* pools WNV+



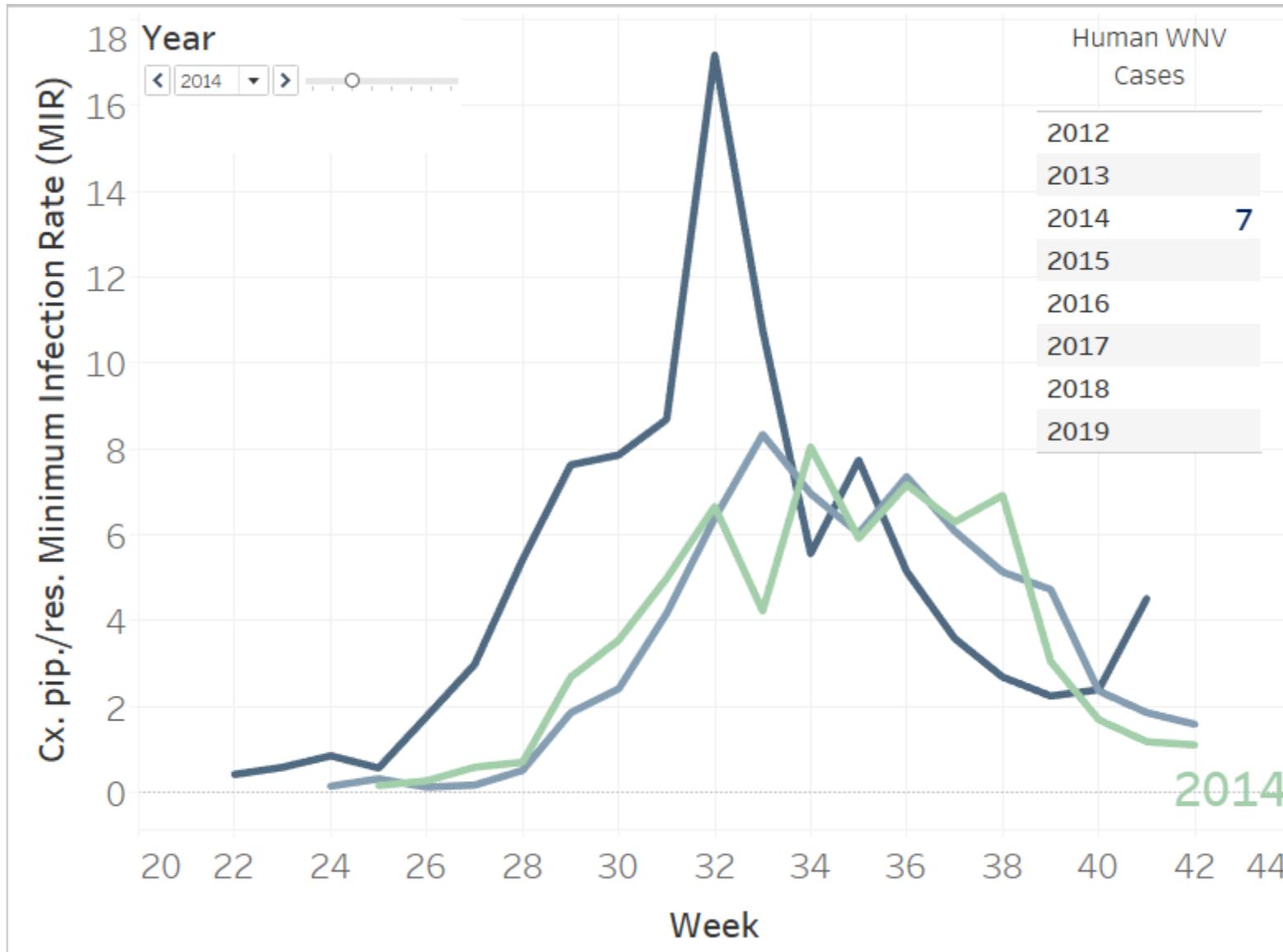
Minimum Infection Rate: $([\text{number of positive pools} / \text{total specimens tested}] \times 1000)$, with the data representing a single species or species group collected over a time period and geographic area relevant to the goals of the surveillance program.- CDC



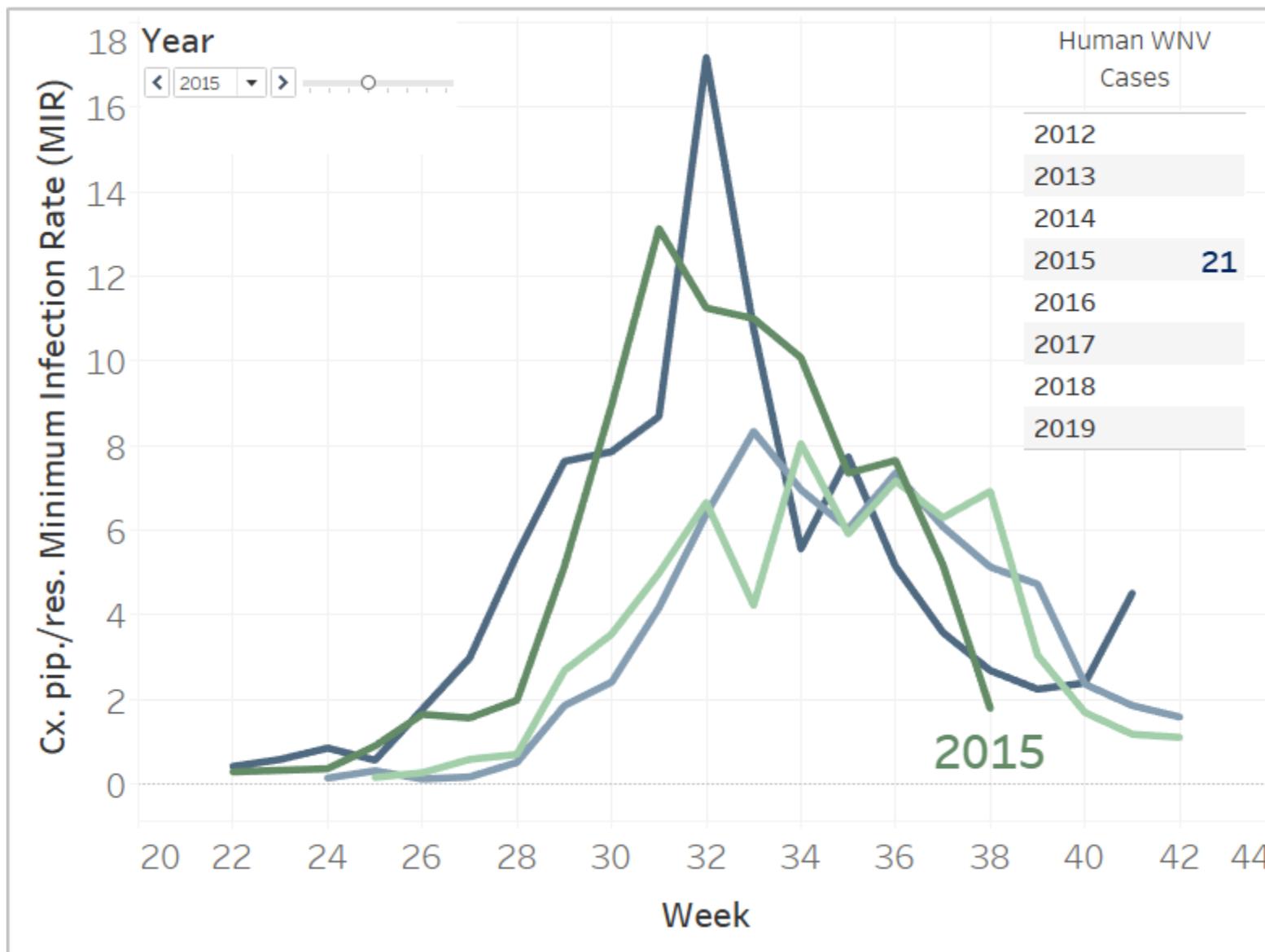
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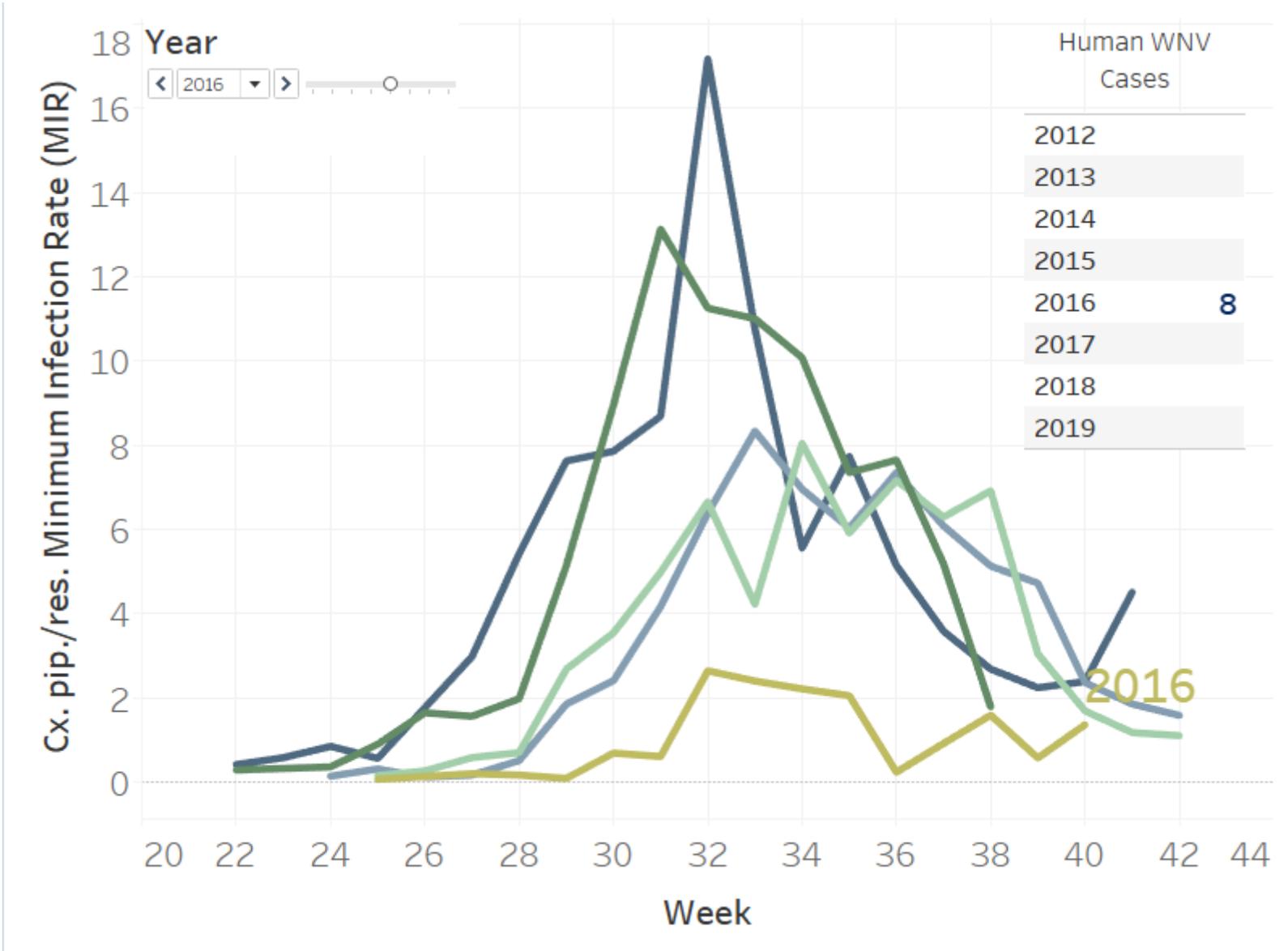
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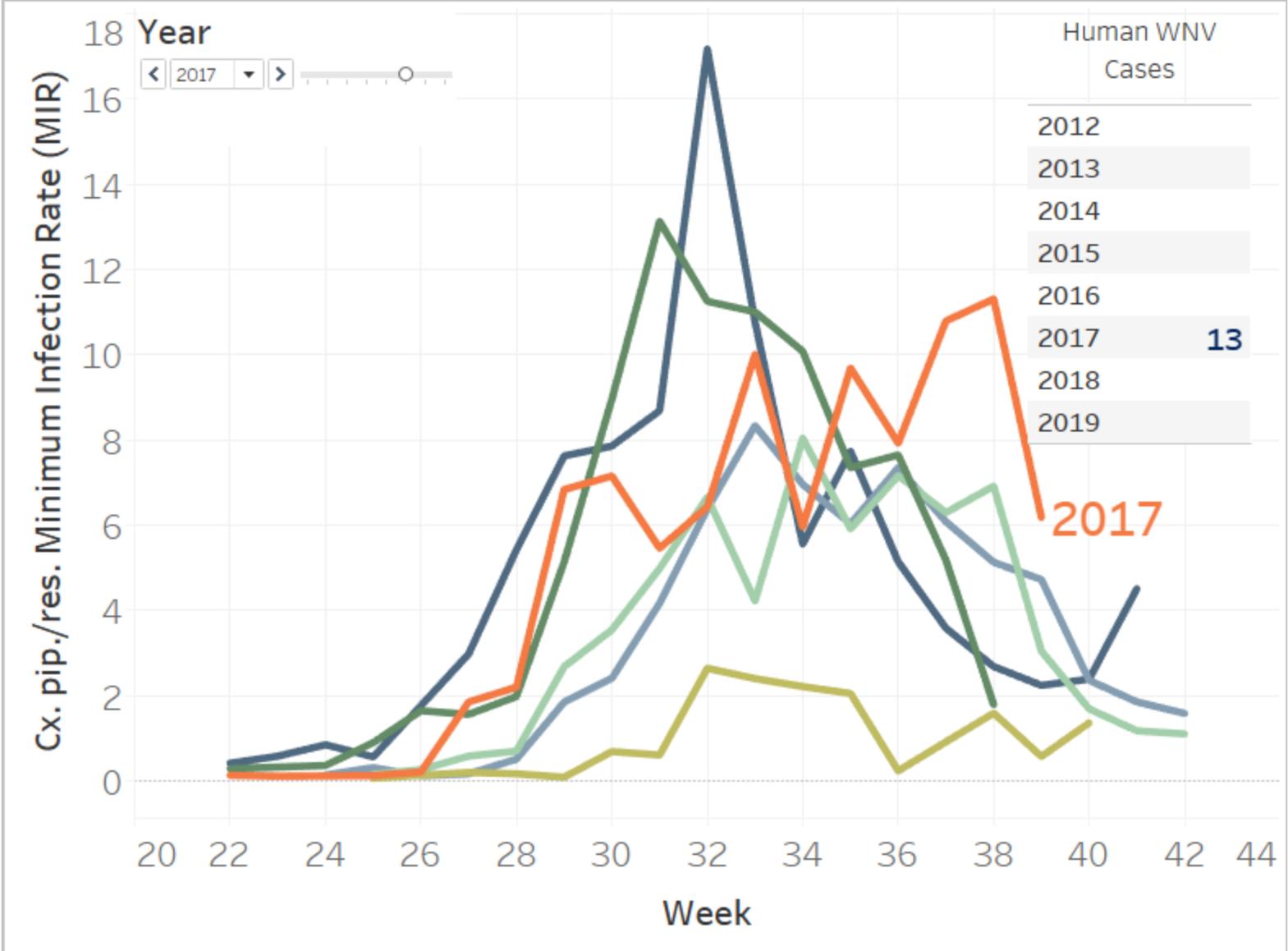
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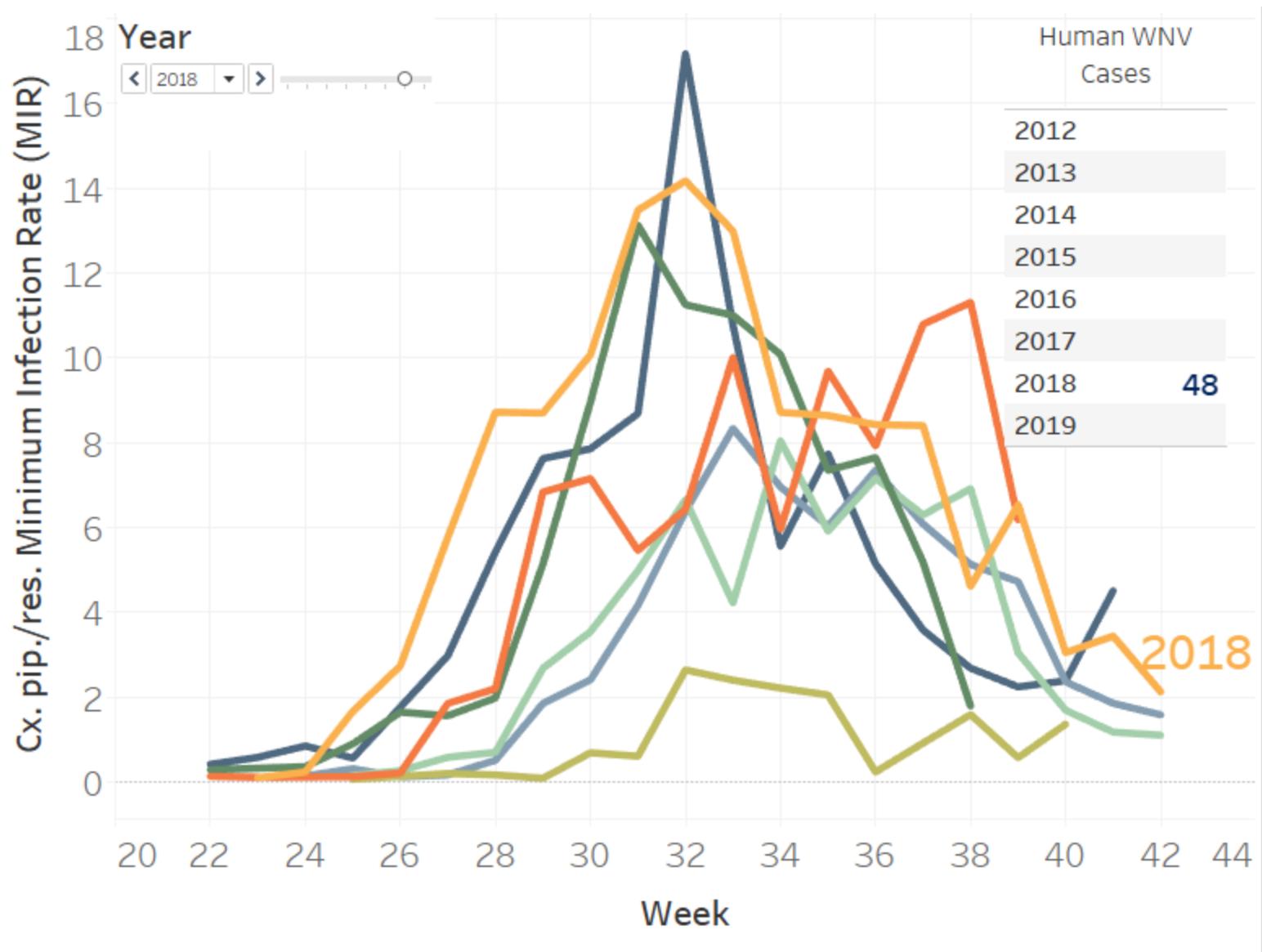
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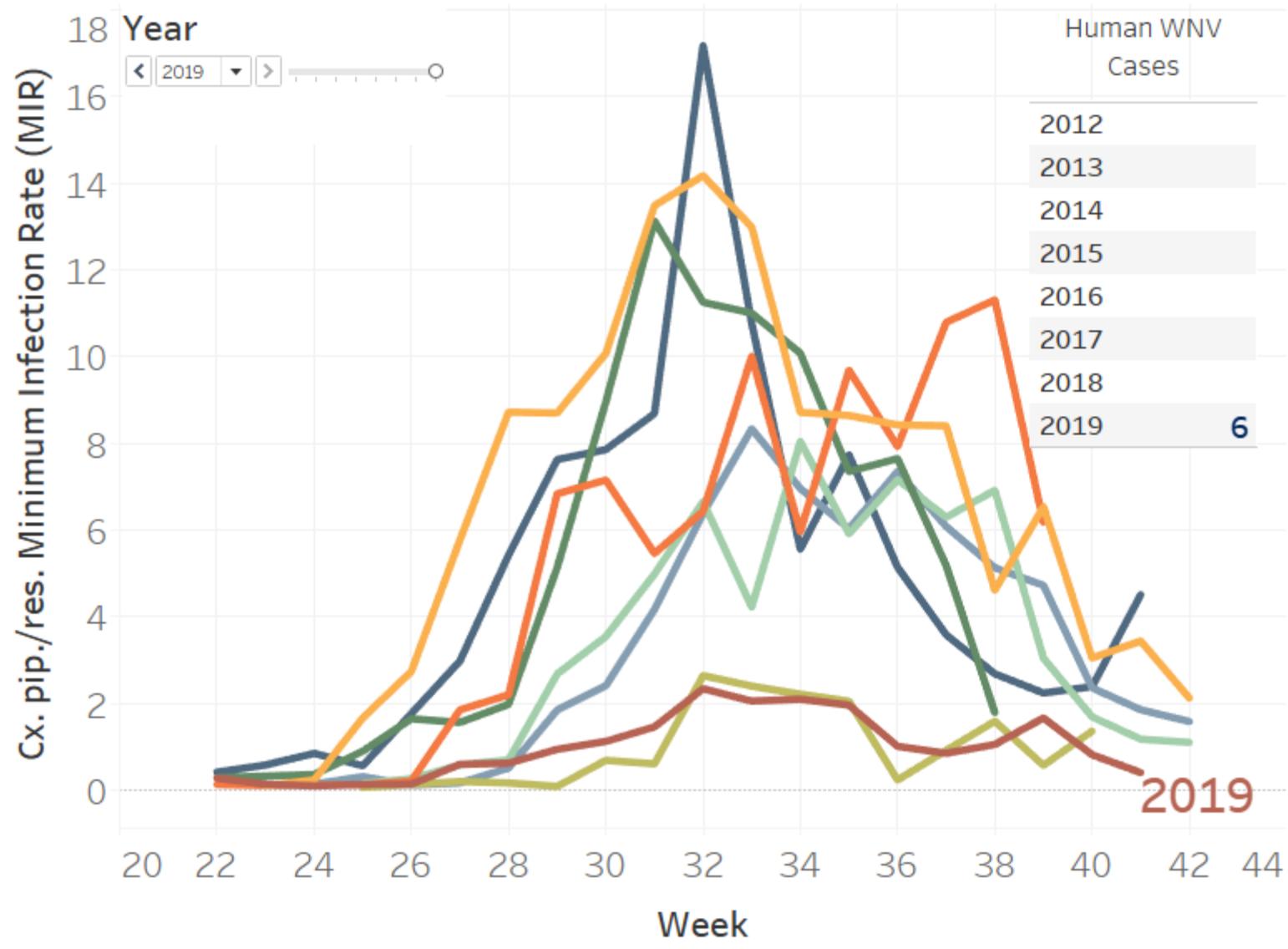
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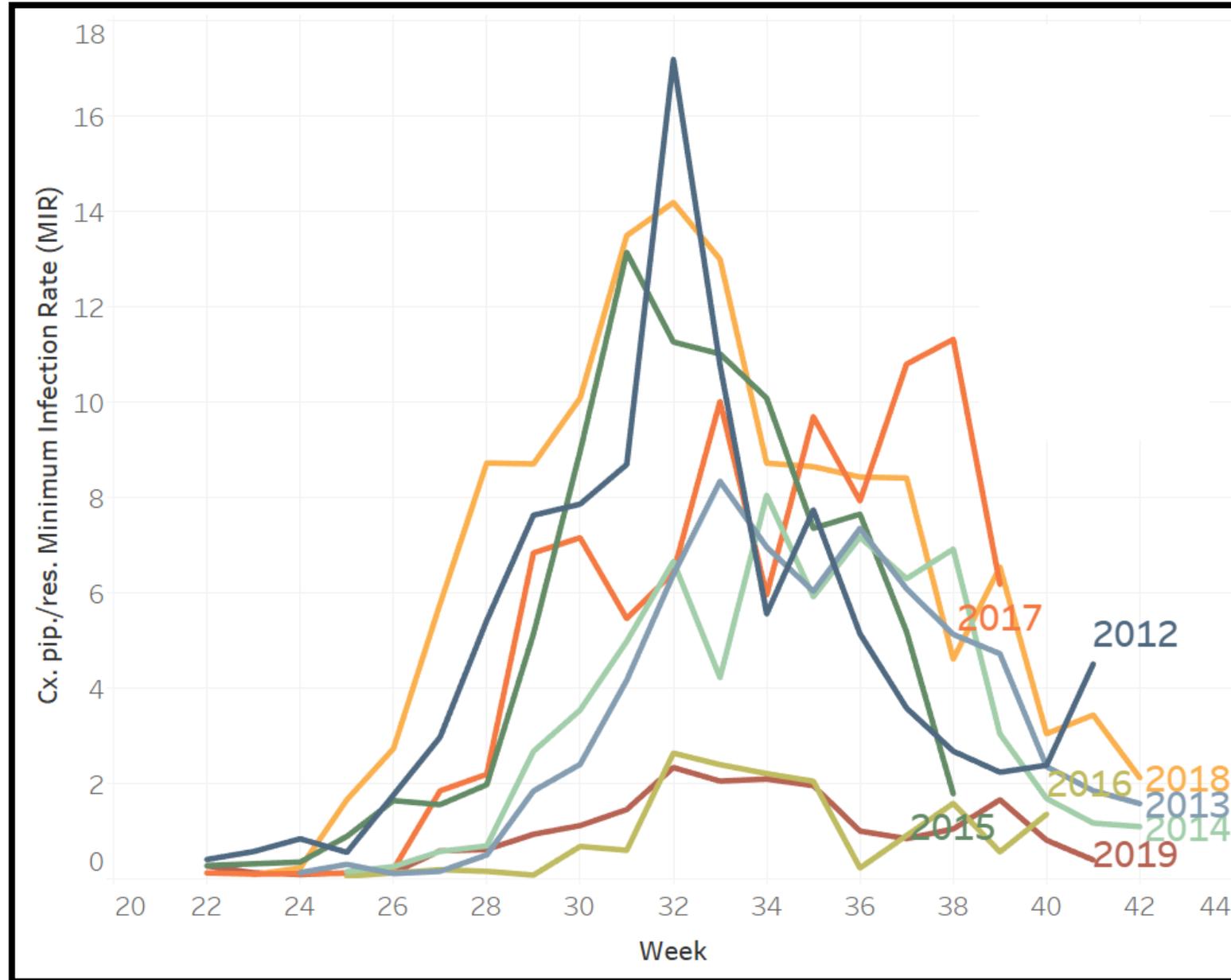


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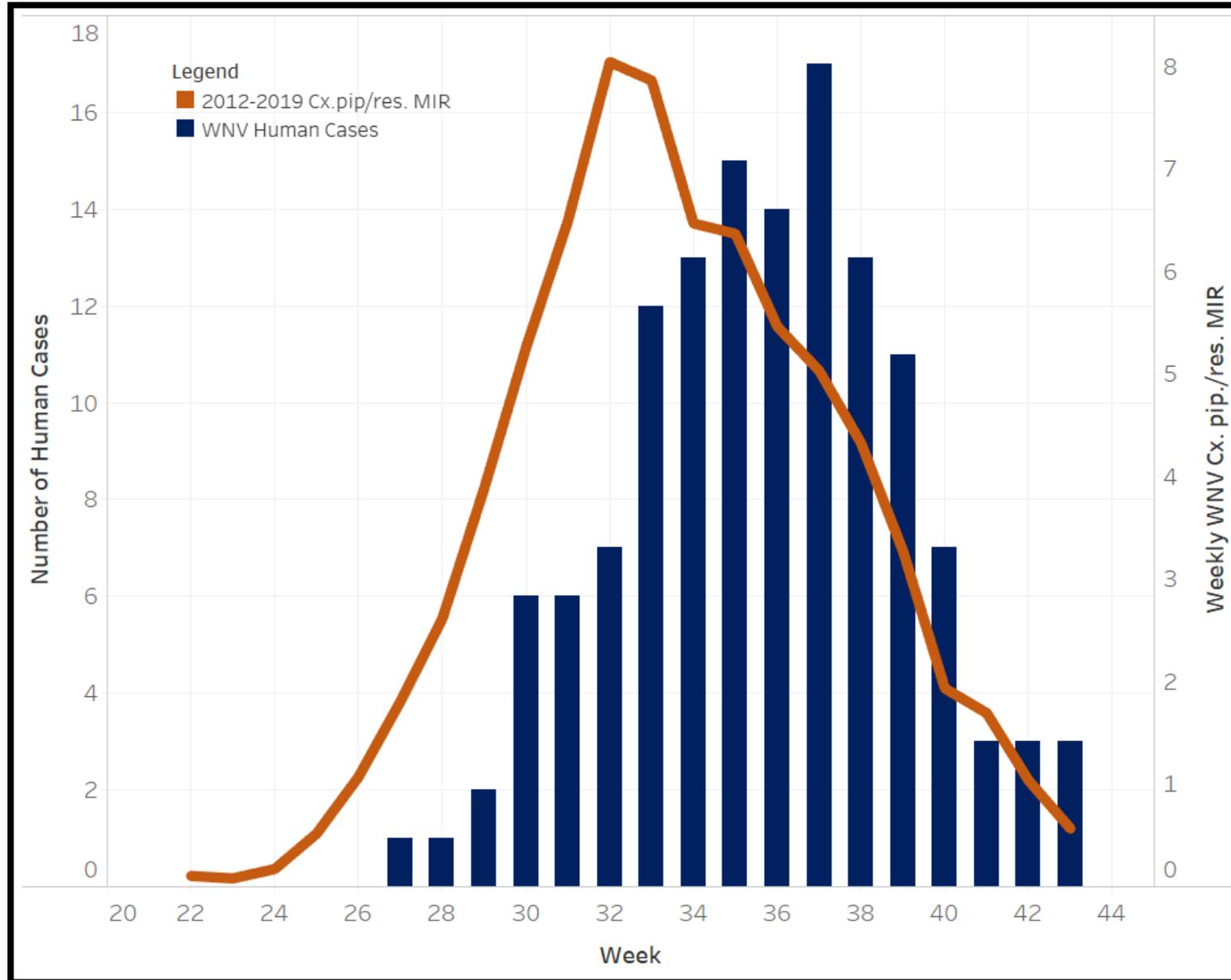
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Weekly Cx. pip./res. MIR 2012-2019

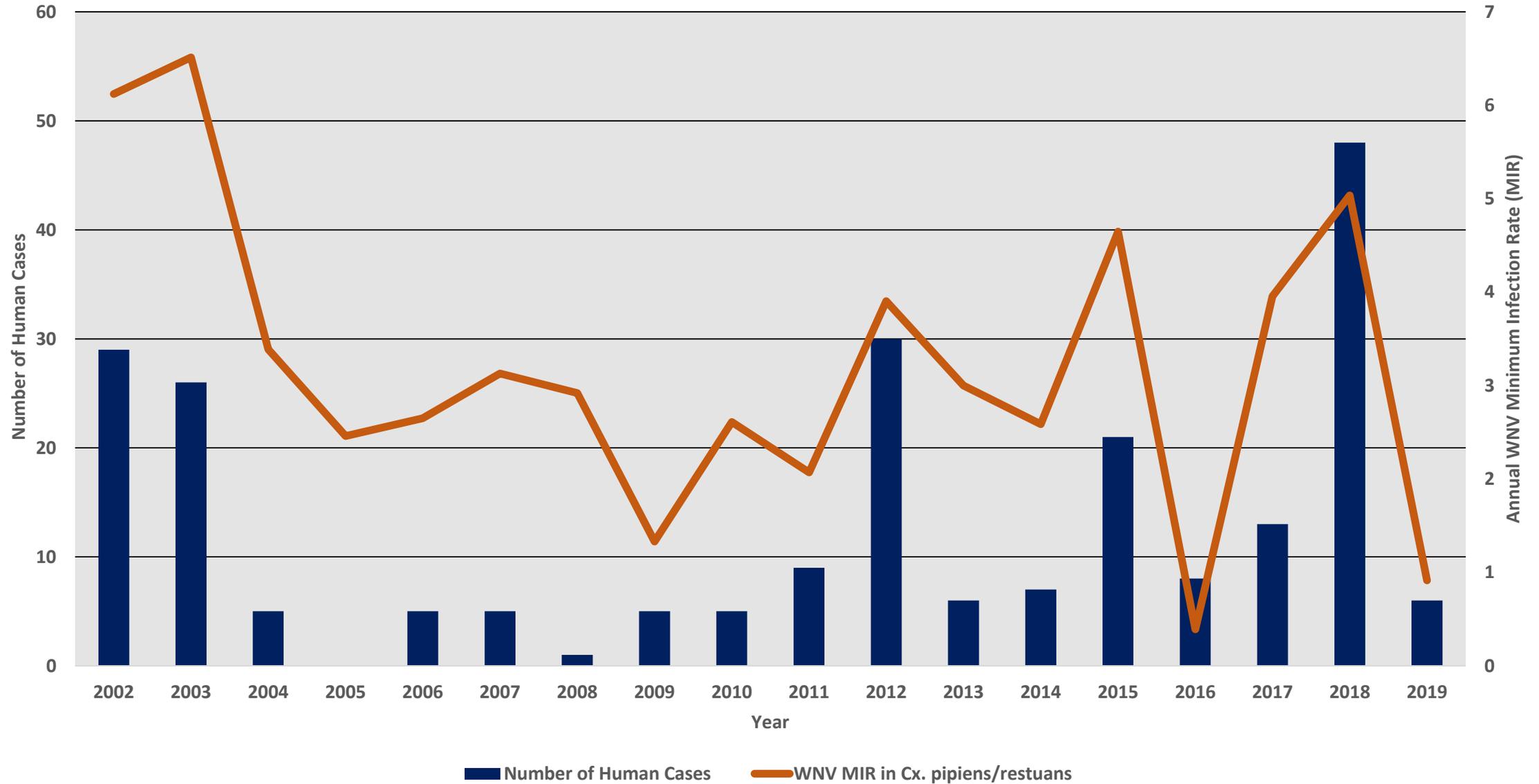


Year	Human Cases
2012	30
2013	6
2014	7
2015	21
2016	8
2017	13
2018	48
2019	6

Weekly Cx. pip./res. MIR 2012-2019



Influence of *Cx. pipiens/restuans* annual WNV minimum infection rate (MIR) on the count human WNV cases in VA.

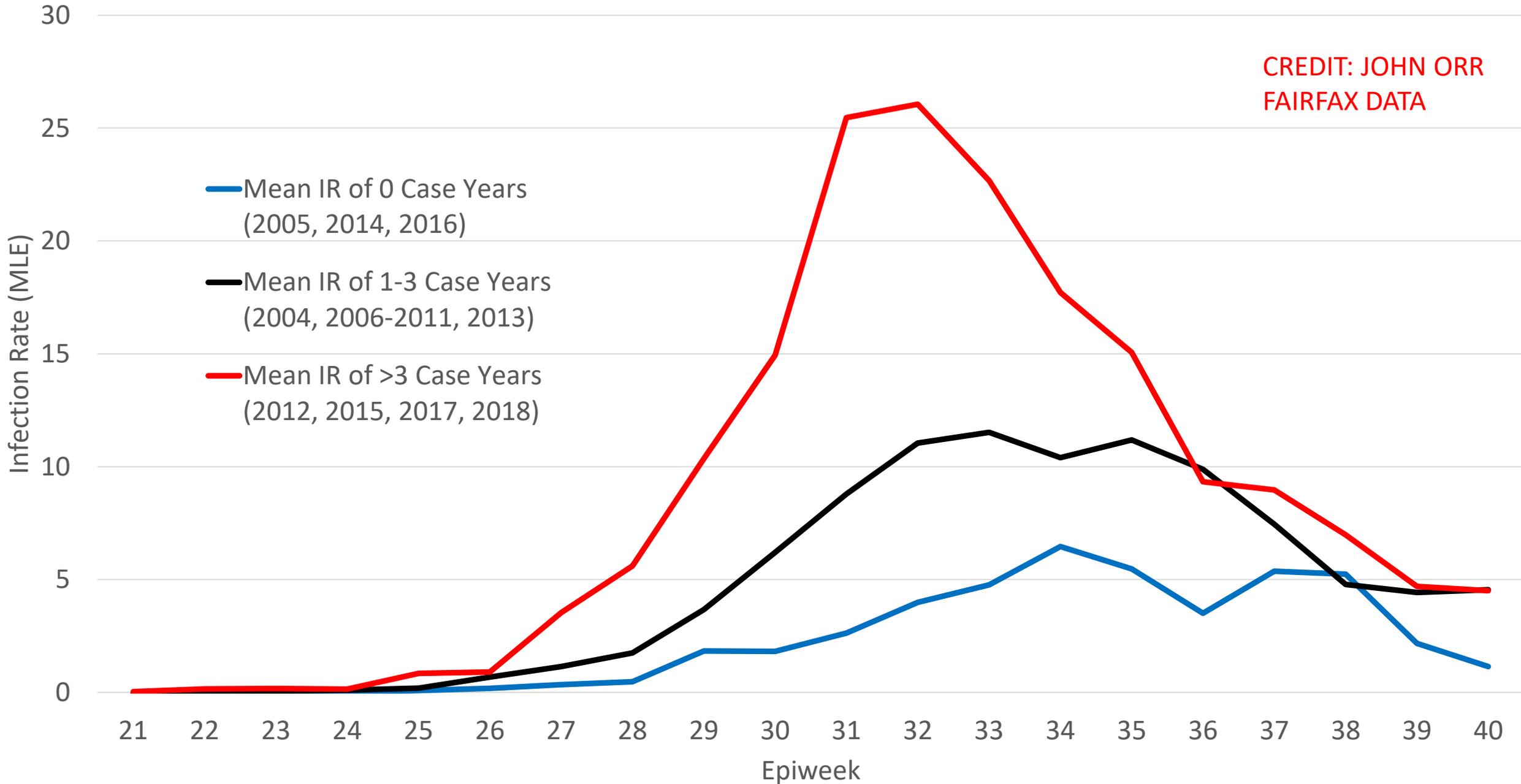


Linking Mosquito and Human WNV Data

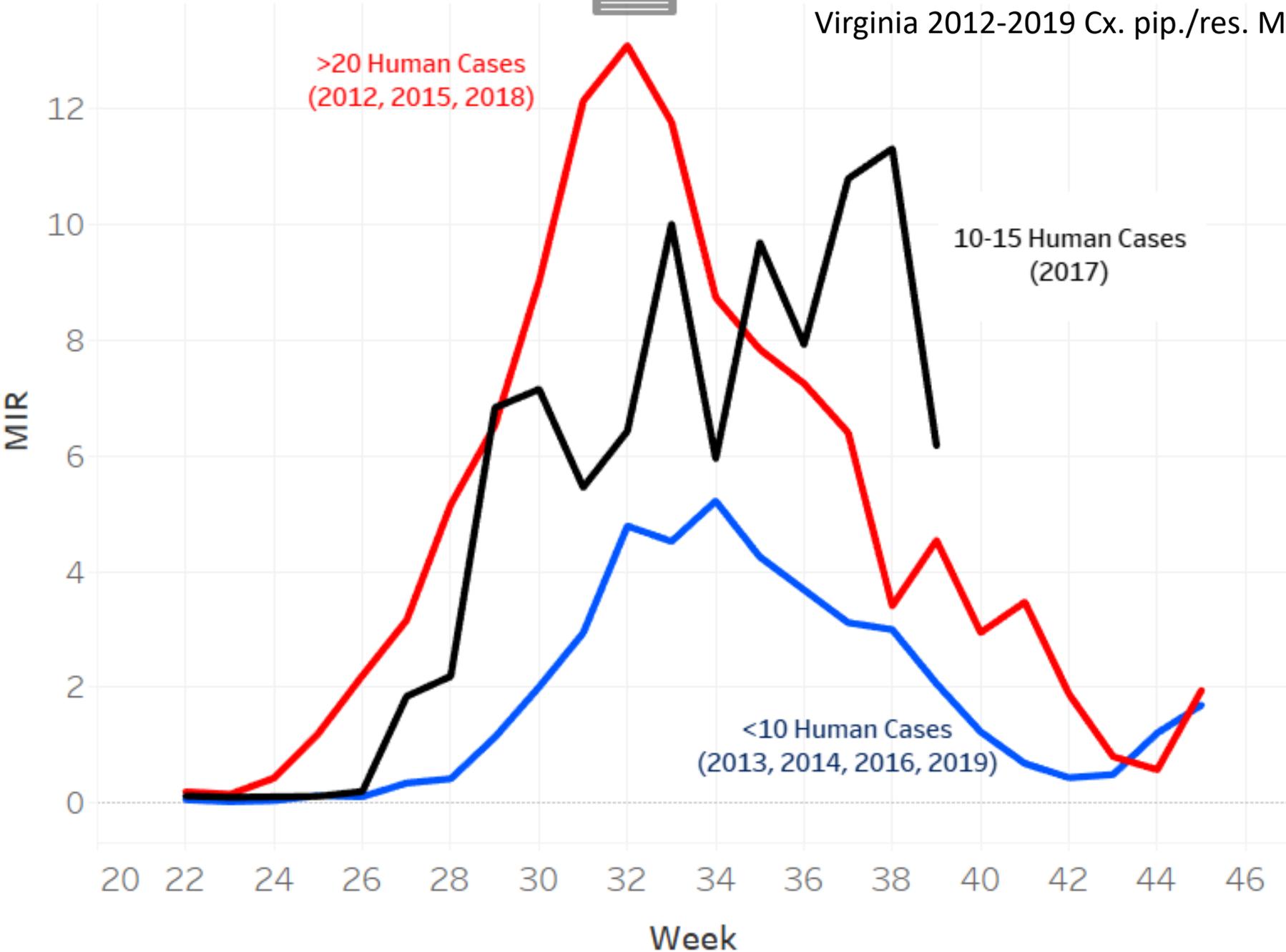
“Holy grail of WNV mosquito surveillance”~John Orr

Culex WNV Infection Rate (MLE), 2004-2018

CREDIT: JOHN ORR
FAIRFAX DATA



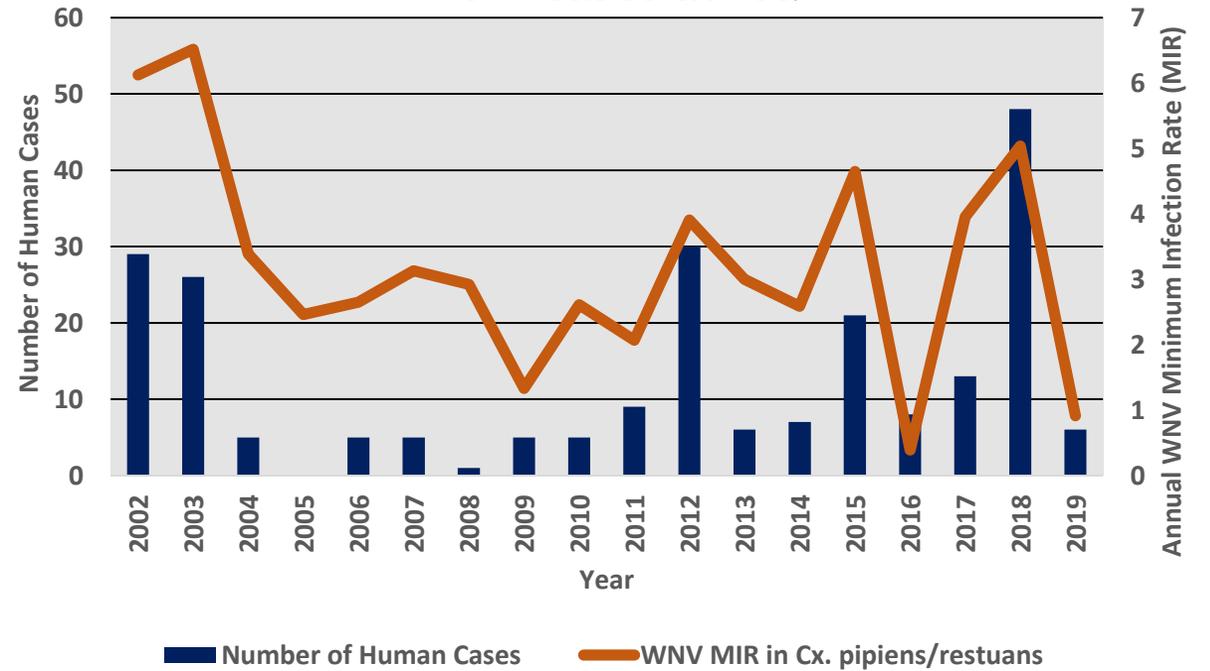
Virginia 2012-2019 Cx. pip./res. MIR



Birds?



Influence of *Cx. pipiens/restuans* annual WNV minimum infection rate (MIR) on the count human WNV cases in VA.



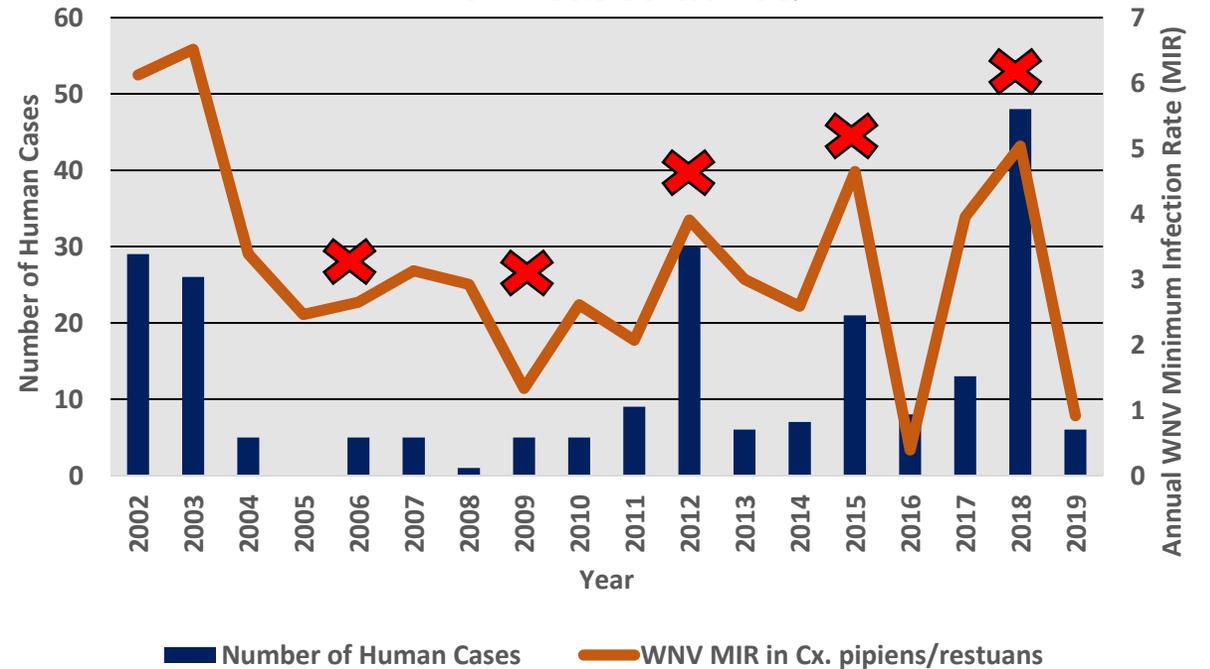
https://www.allaboutbirds.org/guide/American_Robin/overview

<https://www.birdnote.org/show/how-long-does-robin-live>

“An American Robin can produce three successful broods in one year. On average, though, only 40 percent of nests successfully produce young. Only 25 percent of those fledged young survive to November. From that point on, about half of the robins alive in any year will make it to the next. Despite the fact that a lucky robin can live to be 14 years old, the **entire population turns over on average every six years.**”

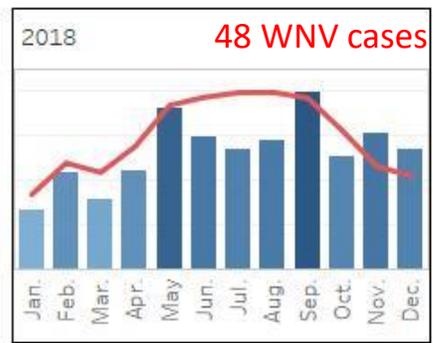
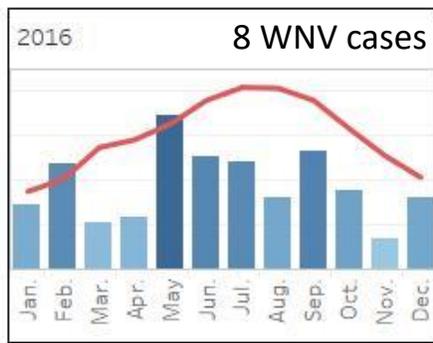
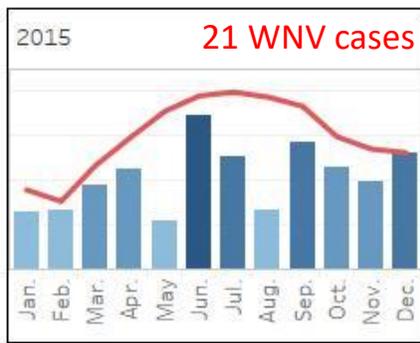
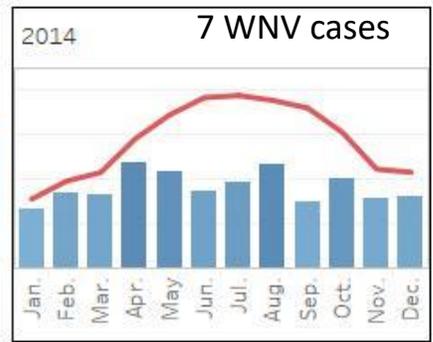
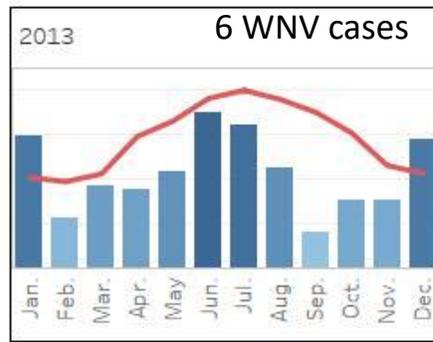
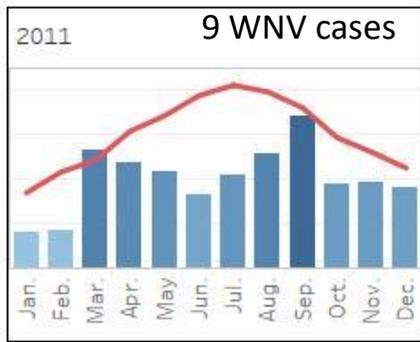
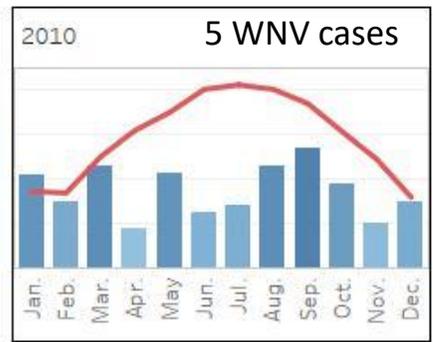
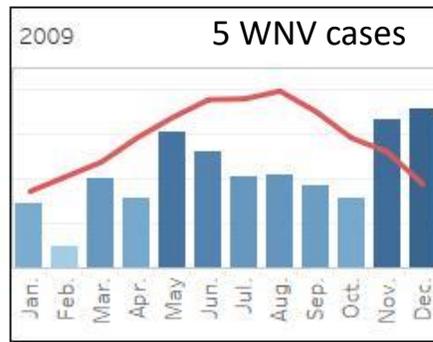
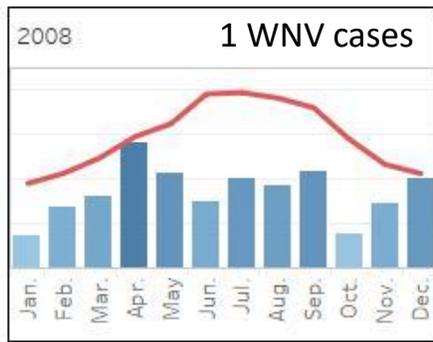
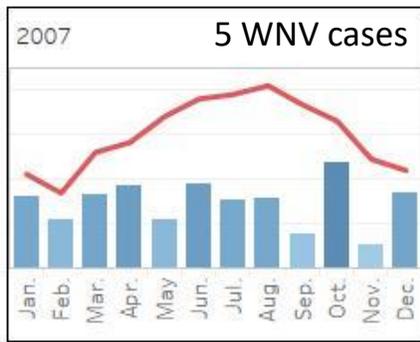
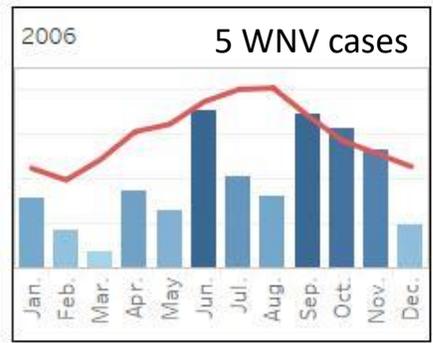
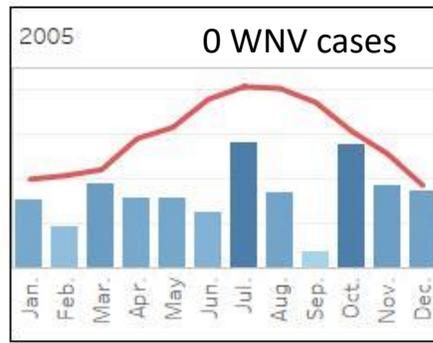
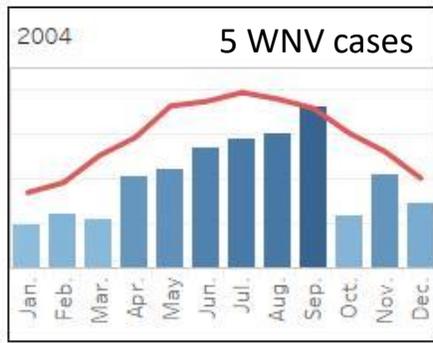
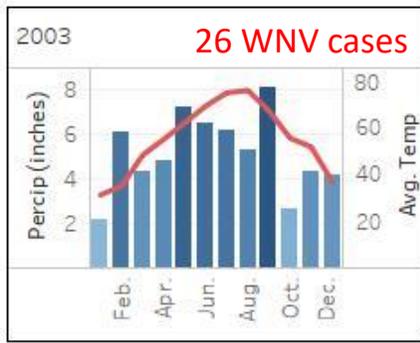


Influence of *Cx. pipiens/restuans* annual WNV minimum infection rate (MIR) on the count human WNV cases in VA.



https://www.allaboutbirds.org/guide/American_Robin/overview

<https://www.birdnote.org/show/how-long-does-robin-live>





Increased transmissibility of Emergent Genotypes of West Nile virus in New York State

Sean M. Bialosuknia¹, Alan P. Dupuis II¹, Yi Tan², Steven D. Zink¹, Cheri A. Koetzner¹, Joseph G. Maffei¹, Rebecca A. Halpin³, Emmi Muller³, Mark Novatny³, Meghan Shilts^{2, 3}, Nadia B. Fedorova³, Paolo Amedeo³, Suman R. Das², Brett Pickett³, Jennifer C. Owen⁴, Hannah Landwerlen⁴, Laura D. Kramer^{1,5} and Alexander T. Ciota^{1,5}

¹The Arbovirus Laboratory, Wadsworth Center, New York State Department of Health, Slingerlands, NY, United States of America

² Department of Medicine, Vanderbilt University Medical Center, Nashville, Tennessee, USA.

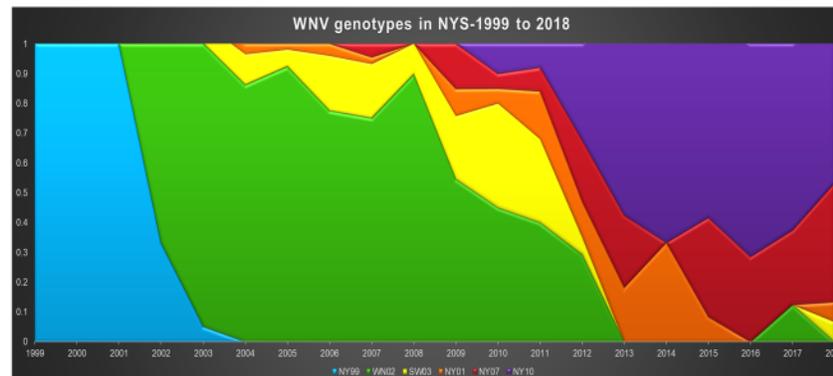
³J. Craig Venter Institute, Rockville, Maryland, USA.

⁴ Department of Fisheries and Wildlife, Michigan State University, East Lansing, MI USA.

⁵Department of Biomedical Sciences, State University of New York at Albany School of Public Health, Albany, NY, United States of America

HIGHLIGHTS

- Phylogenetic and selection analyses of over 1200 full-genome WNV sequences, focusing on ~500 newly sequenced NYS WNV isolates from 1999-2018
- Identified 13 positions in the WNV genome under positive selection in the US, including 7 in NYS
- Novel genotypes with shared, positively selected sites have emerged in NYS over the last decade: NY07 and NY10
- Genotype displacement was concurrent with increased WNV activity from 2010-18
- NY10 strains are more infectious and transmissible by *Cx. pipiens* and American robins, supporting the hypothesis that adaptive evolution drove WNV displacement and contributed to increased WNV activity



Department of Health

Wadsworth Center

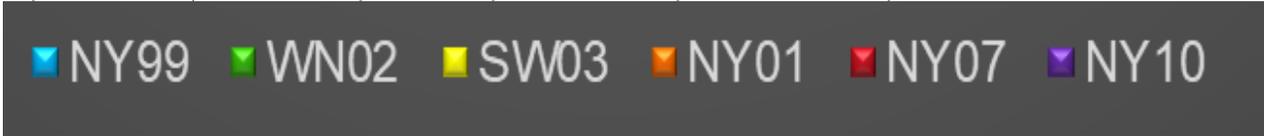
Increased transmissibility of Emergent Genotypes of West Nile virus in New York State

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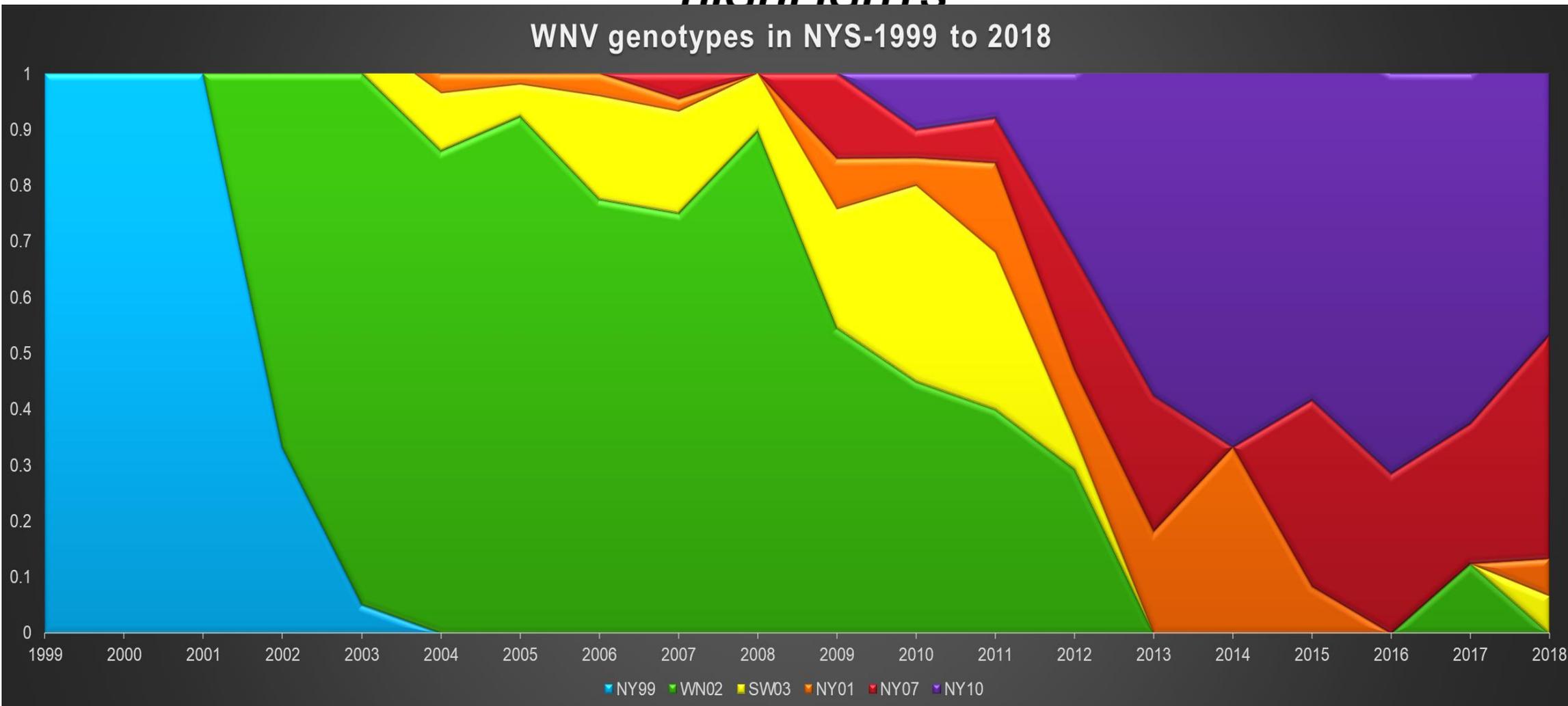
NEVBD
NORTHEAST REGIONAL CENTER FOR
EXCELLENCE IN VECTOR-BORNE DISEASES

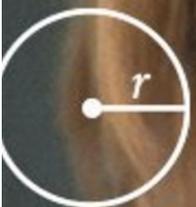
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HIGHLIGHTS

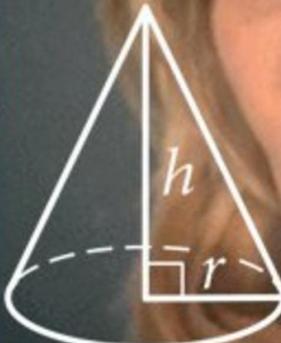
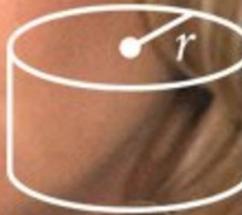
WNV genotypes in NYS-1999 to 2018





$A = \pi r^2$
 $C = 2\pi r$

$V = \frac{1}{3} \pi r^2 h$

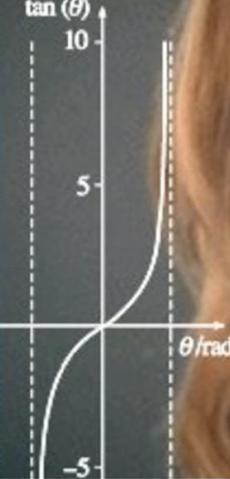



$V = \pi r^2 h$

	30°	45°	60°
sin	$\frac{1}{2}$	$\frac{\sqrt{2}}{2}$	$\frac{\sqrt{3}}{2}$
cos	$\frac{\sqrt{3}}{2}$	$\frac{\sqrt{2}}{2}$	$\frac{1}{2}$
tan	$\frac{\sqrt{3}}{3}$	1	$\sqrt{3}$

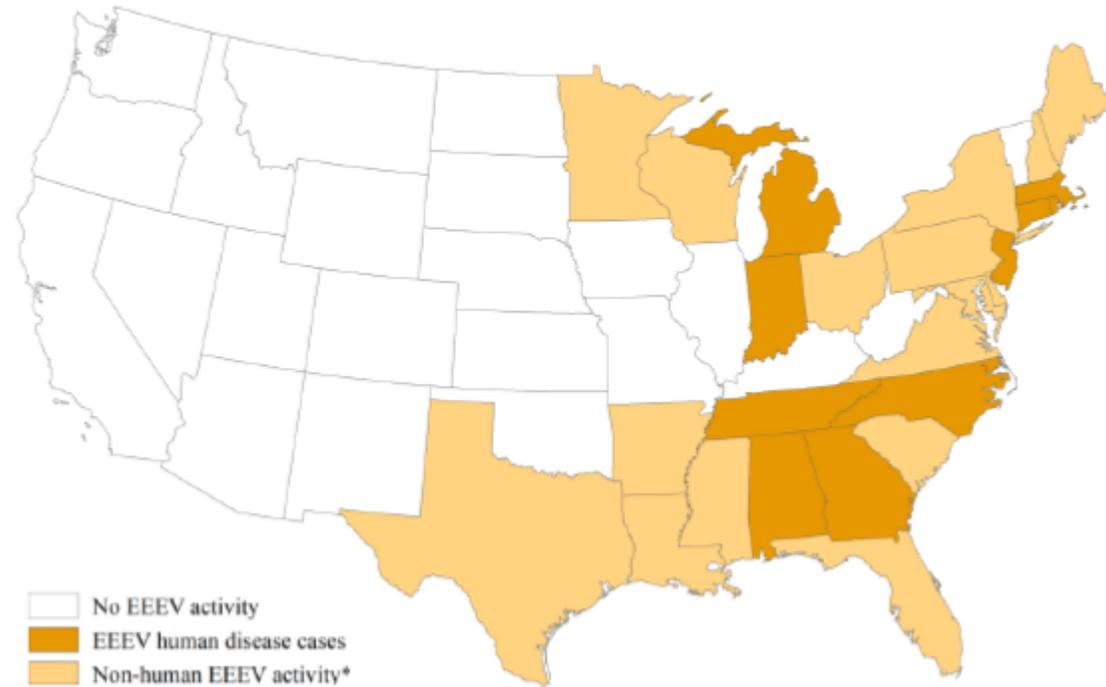



$\int \sin x dx = -\cos x + C$
 $\int \frac{dx}{\cos^2 x} = \tan x + C$
 $\int \tan x dx = -\ln|\cos x| + C$
 $\int \frac{dx}{\sin x} = \ln\left|\frac{x}{2}\right| + C$
 $\int \frac{dx}{a^2 + x^2} = \frac{1}{a} \arctg \frac{x}{a} + C$
 $\int \frac{dx}{x^2 - a^2} = \frac{1}{2a} \ln\left|\frac{x-a}{x+a}\right| + C$



$ax^2 + bx + c = 0$
 $a\left(x^2 + \frac{b}{a}x + \frac{c}{a}\right) = 0$
 $x^2 + 2\frac{b}{2a}x + \left(\frac{b}{2a}\right)^2 - \left(\frac{b}{2a}\right)^2 + \frac{4ac}{4a^2} = 0$
 $\left(x + \frac{b}{2a}\right)^2 - \frac{b^2 - 4ac}{4a^2} = 0$

Figure 4. Eastern equine encephalitis virus (EEEV) activity reported to ArboNET, by state — United States, 2019 (as of January 7, 2020)



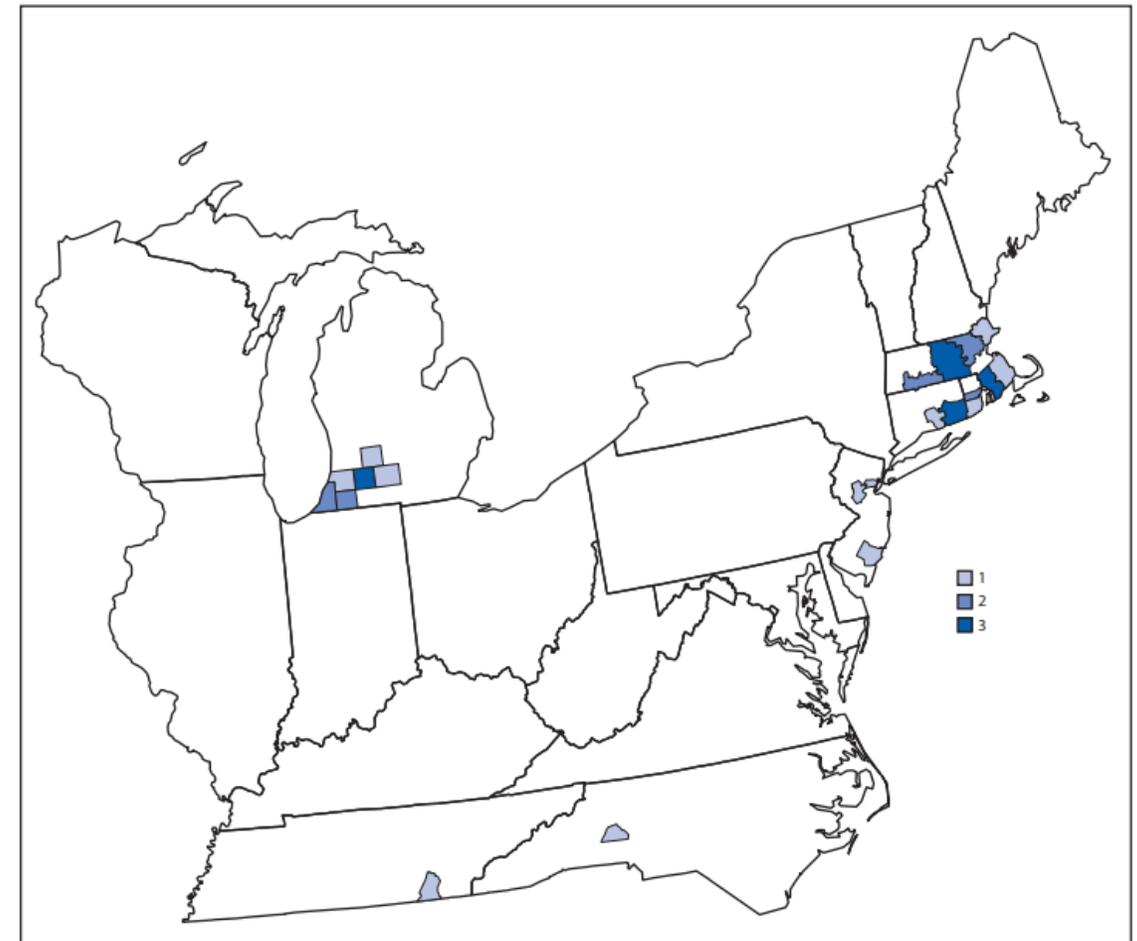
*EEEV veterinary disease cases, or infections in mosquitoes, birds, or sentinel animals

Table 2. Eastern equine encephalitis virus human disease cases reported to ArboNET, United States, 2019

State	Neuroinvasive disease cases	Non-neuroinvasive disease cases	Total cases*	Deaths
Alabama	1	0	1	1
Connecticut	4	0	4	3
Georgia	1	0	1	0
Indiana	1	0	1	1
Massachusetts	12	0	12	3
Michigan	10	0	10	6
New Jersey	4	0	4	0
North Carolina	1	0	1	0
Rhode Island	3	0	3	1
Tennessee	1	0	1	0
Totals	38	0	38	15

*Includes confirmed and probable cases.

FIGURE. Number of reported cases of Eastern equine encephalitis virus disease (N = 34), by county of residence — United States, 2019*

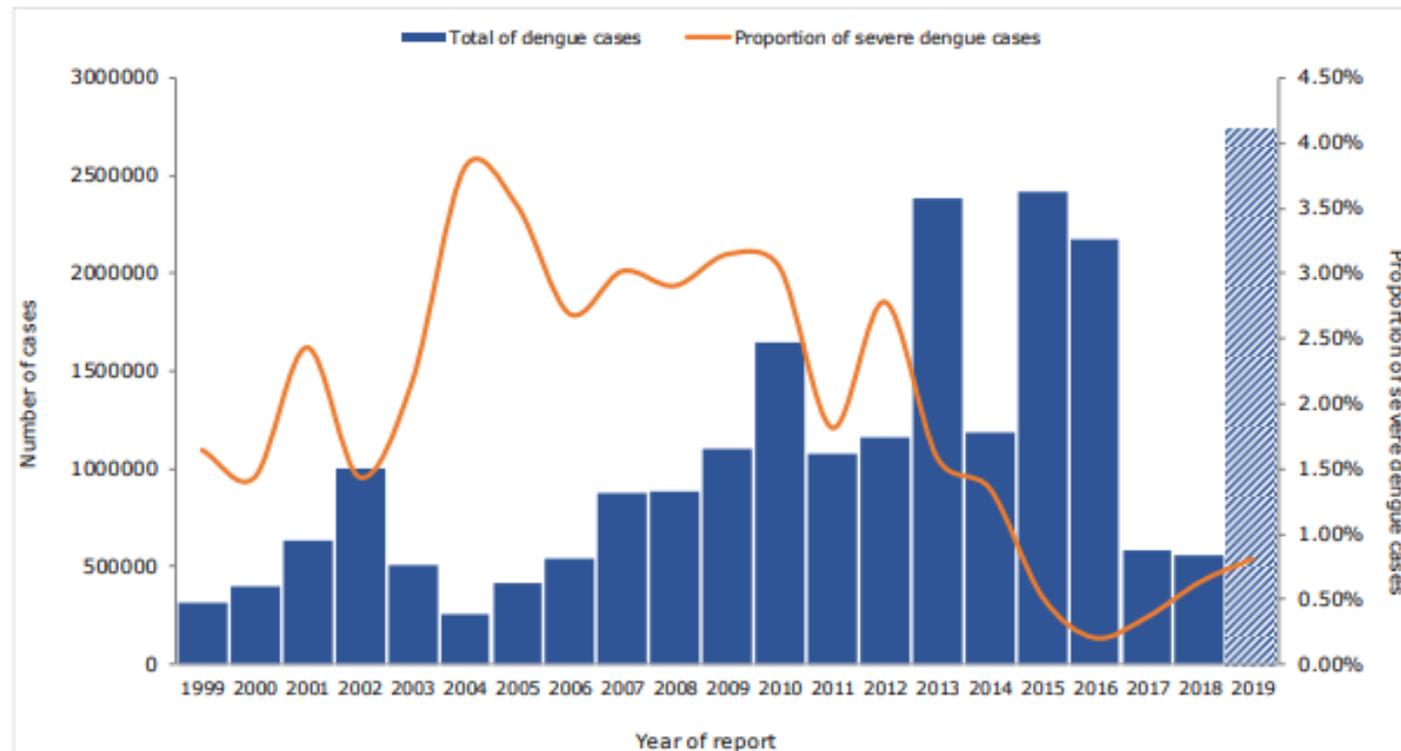


Lindsey N. et al. Multistate Outbreak of Eastern Equine Encephalitis Virus—United States, 2019 (2020) MMWR

Dengue

“Rapid unplanned urbanization, changing land use patterns and increased international travel and trade have brought people into more frequent contact with vectors; climate and other environmental changes have added to their spread.” - PAHO

Figure 1. Distribution of reported dengue cases and proportion of severe dengue cases, by year of report. Region of the Americas, 1999-2019 (up to EW 42 of 2019).



Source: Data entered into the Health Information Platform for the Americas (PLISA, PAHO / WHO) by the Ministries and Institutes of Health of the countries and territories of the Region.

Accomplishments/Reminders

- Mosquito Monthly Report (<http://www.vdh.virginia.gov/environmental-epidemiology/bugs-human-health-statistics/>)
- Mosquito data is available on ArboNET for current year
- Attending TRAST, NMMM
- No mosquito testing at DCLS for 2020
- Enter IR data to MosquitoNET PLZ
- NEVBD CoE Newsletter interest

Questions?

MANY THANKS TO:

- West Nile Virus Northern VA Managers Meeting
- TRAST
- VMCA
- Chesapeake Mosquito Control
- Fairfax County Dept. of Health- Vector Borne Diseases Program
- Hampton Mosquito Control
- Henrico County Mosquito Control
- Norfolk Dept. of Health Vector Control
- Suffolk Mosquito Control
- Virginia Beach Mosquito Control
- Portsmouth Mosquito Control
- York County Mosquito Control
- Prince William County Vector Control
- The Virginia Division of Consolidated Laboratory Services (DCLS)
- Fairfax County Dept. of Health Laboratory
- The Virginia Dept. of Agriculture and Consumer Services (VDACS) Veterinary Laboratories
- Alex Ciota & New York Health Dept. Wadsworth Lab
- VDH- Vector-borne Team
- VDH- Regional and District Epidemiology Staff



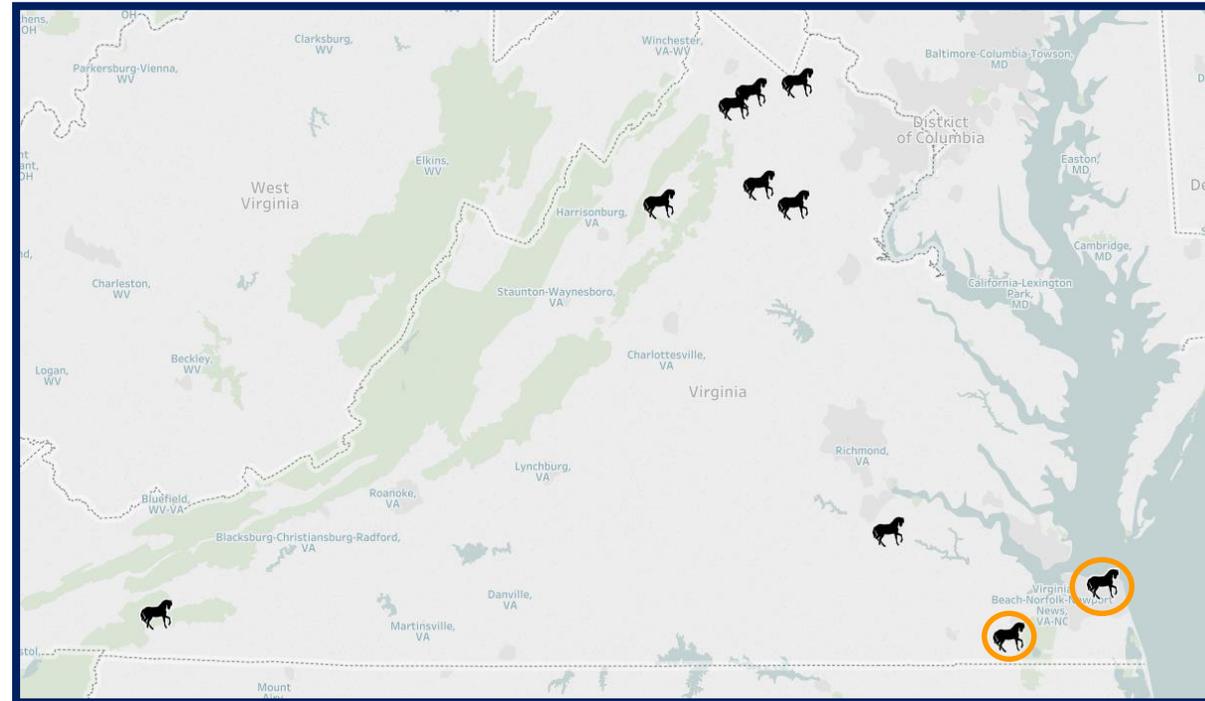
Veterinary Arboviral in VA in 2018

8 reported WNV equine cases

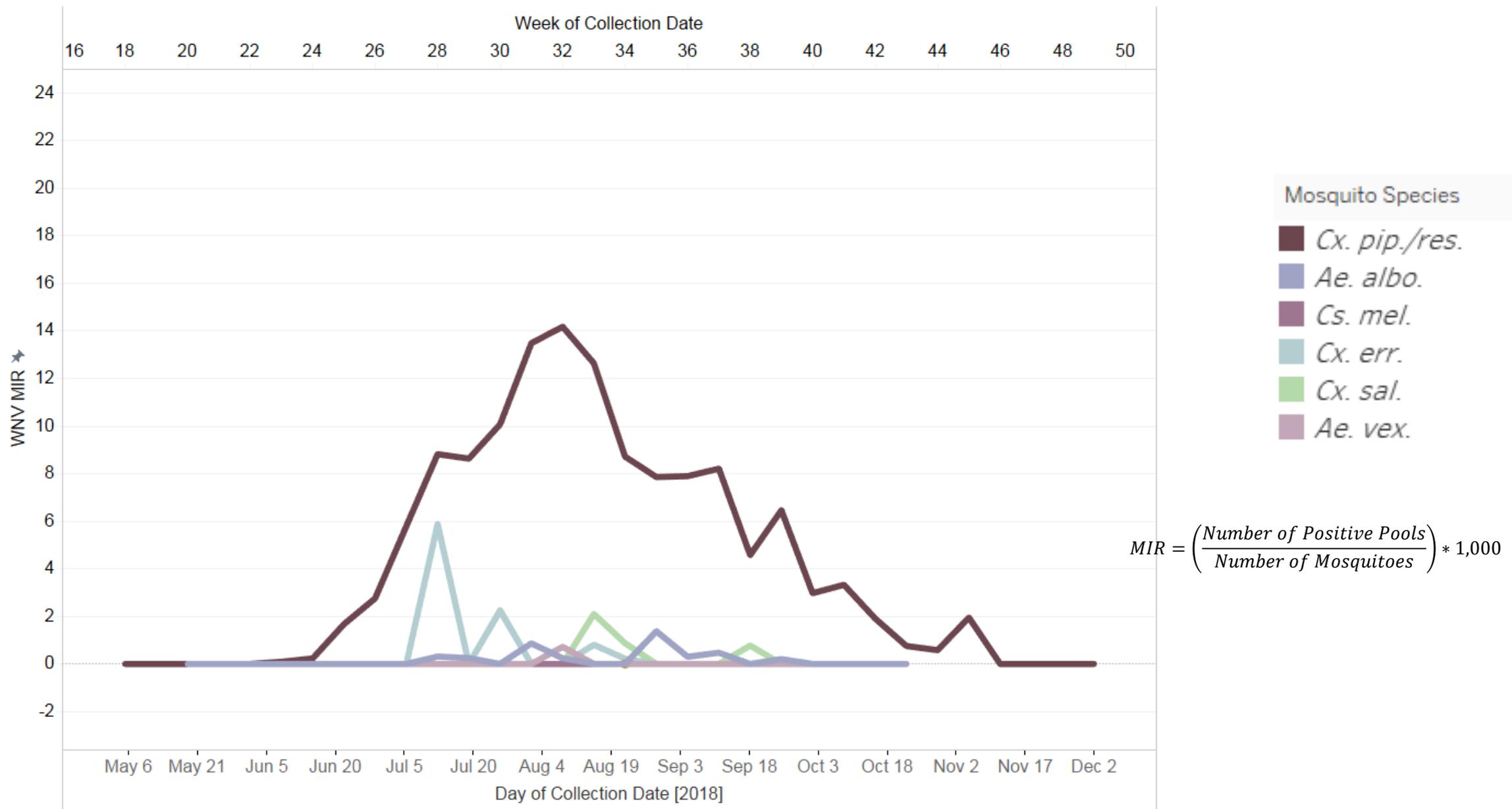
- Fauquier, Warren, Prince George, Page, Loudoun, Rappahannock, Smyth, Clarke

2 reported **EEE** equine cases

- Suffolk, Virginia Beach

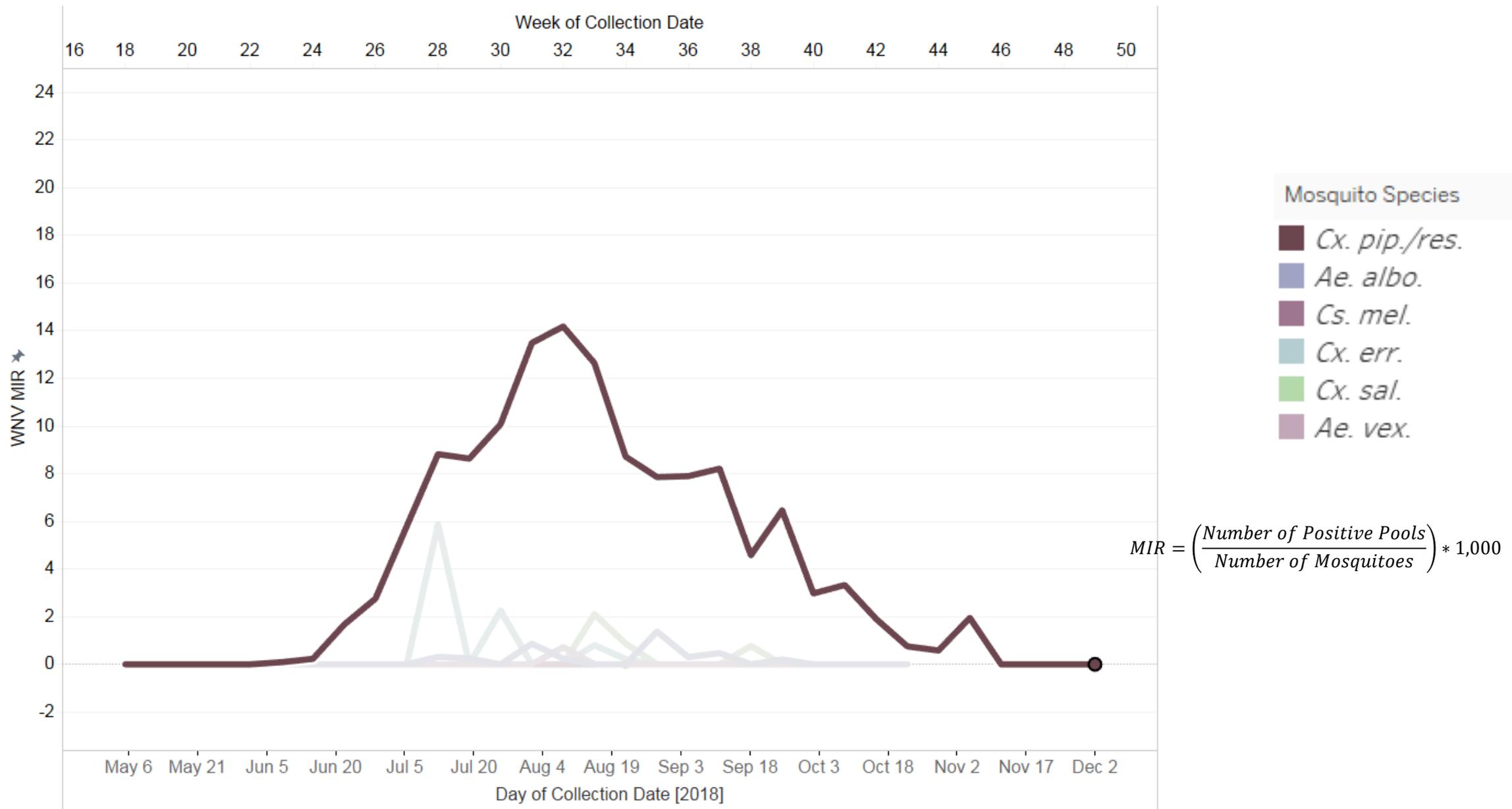


Time series species-specific WNV MIR from May 1-Dec 4.



*Only includes fully identified mosquito species with ≥1,000 mosquitoes trapped/tested (excludes *Ae. tris.*, *Ae. jap.*, *Ae. atl.*, *Cq. per.*, *Cx. spp.*)

Cx. pip./res. WNV MIR peaked at 14.18 MIR during week 32 (Aug 5th-12th).

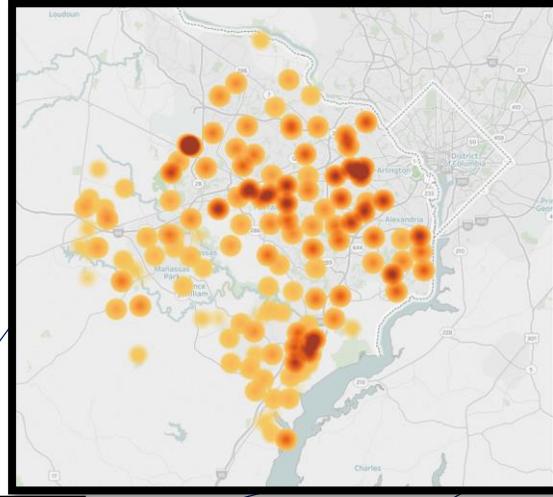


$$MIR = \left(\frac{\text{Number of Positive Pools}}{\text{Number of Mosquitoes}} \right) * 1,000$$

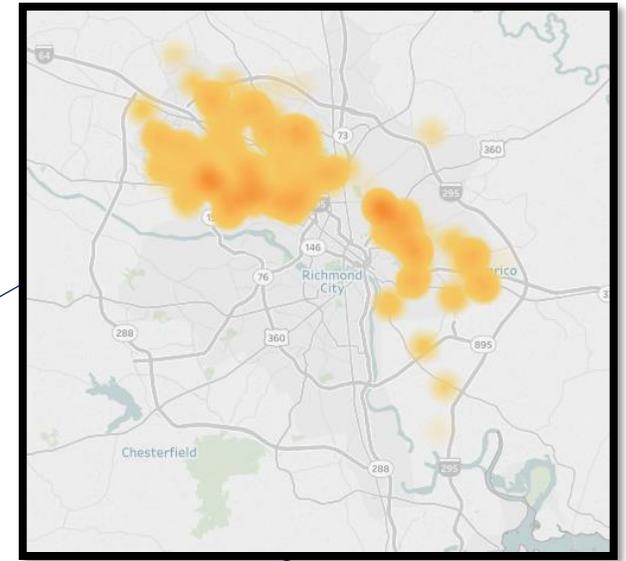
*Only includes fully identified mosquito species with ≥1,000 mosquitoes trapped/tested (excludes *Ae. tris.*, *Ae. jap.*, *Ae. atl.*, *Cq. per.*, *Cx. spp.*)

There were **316,021** mosquitoes trapped and tested in **9,166** pools during 2019 surveillance.

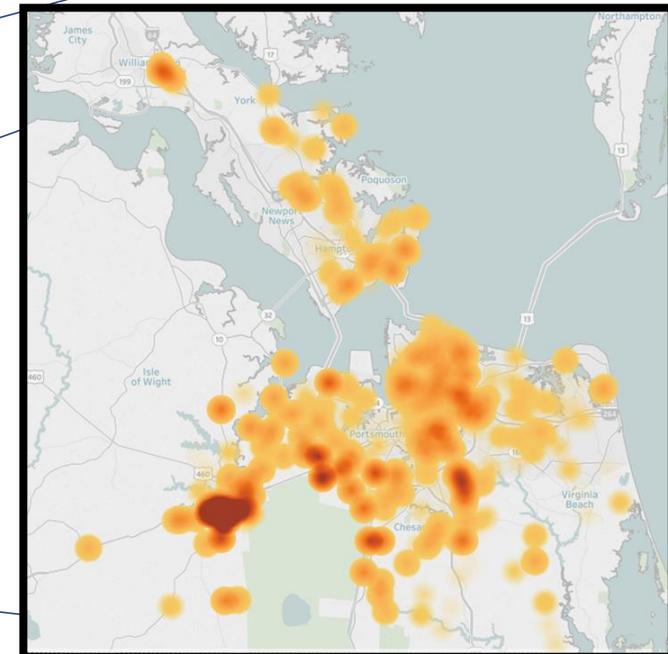
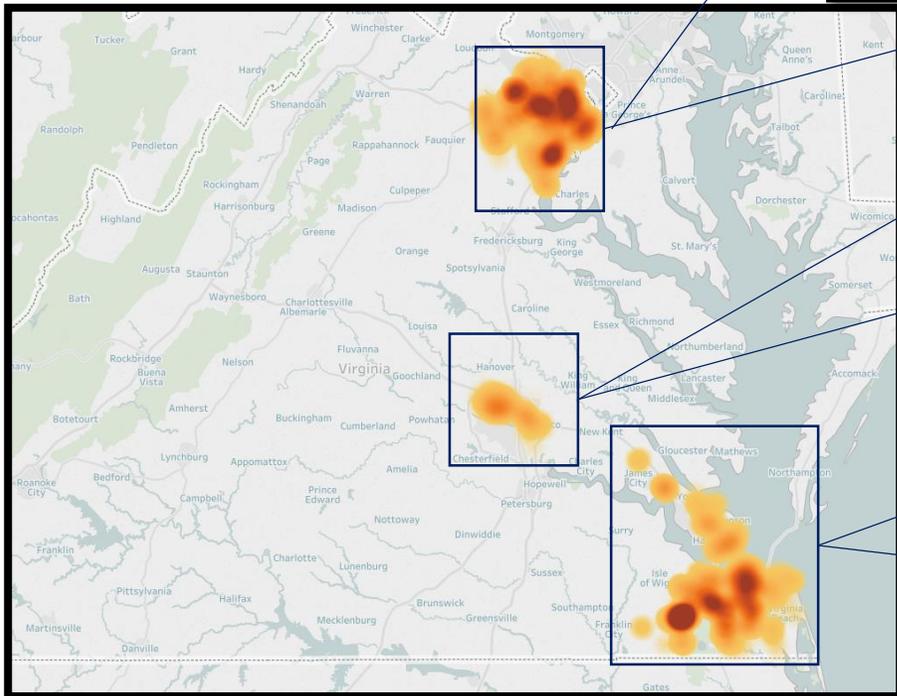
NOVA



Central

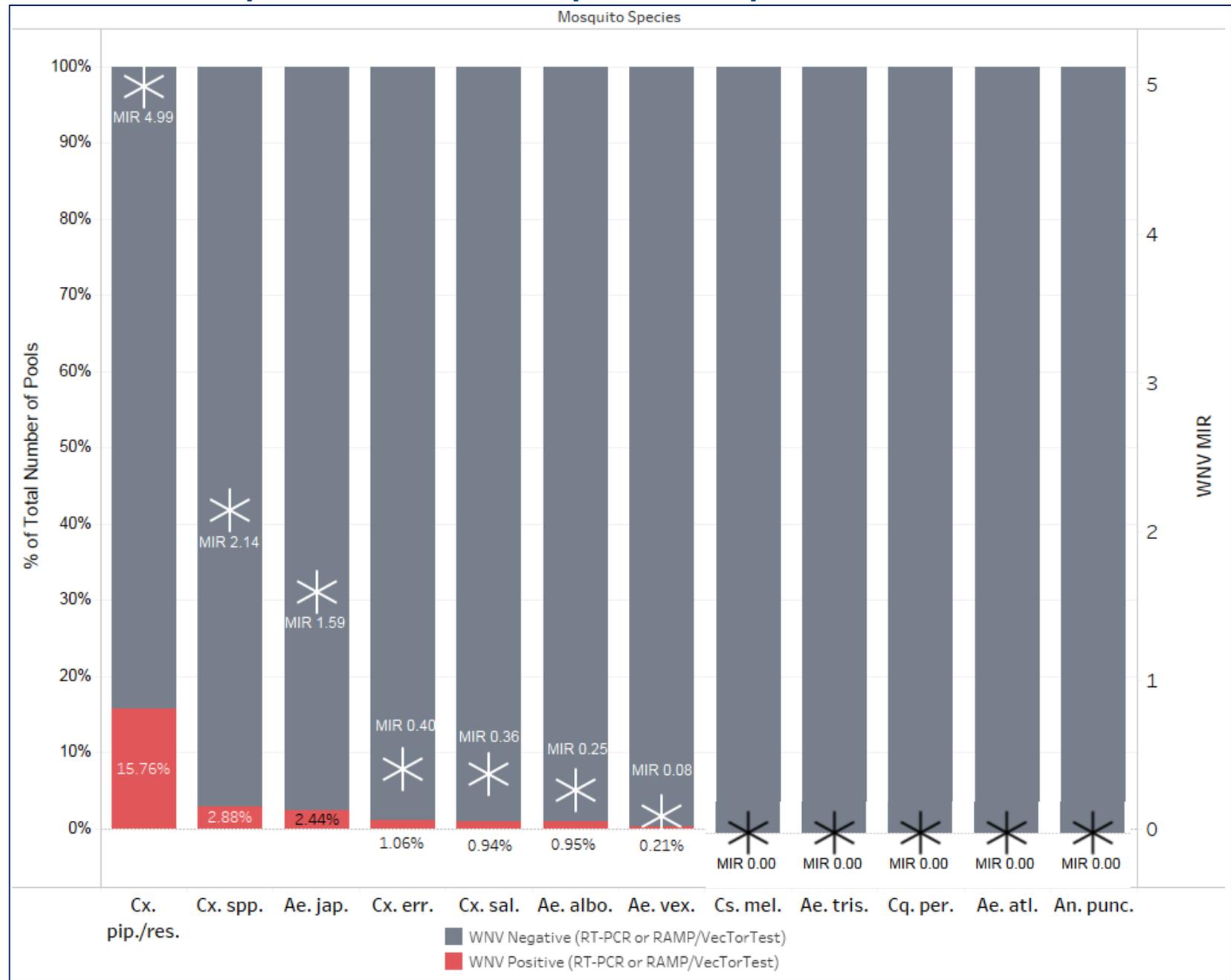


Tidewater



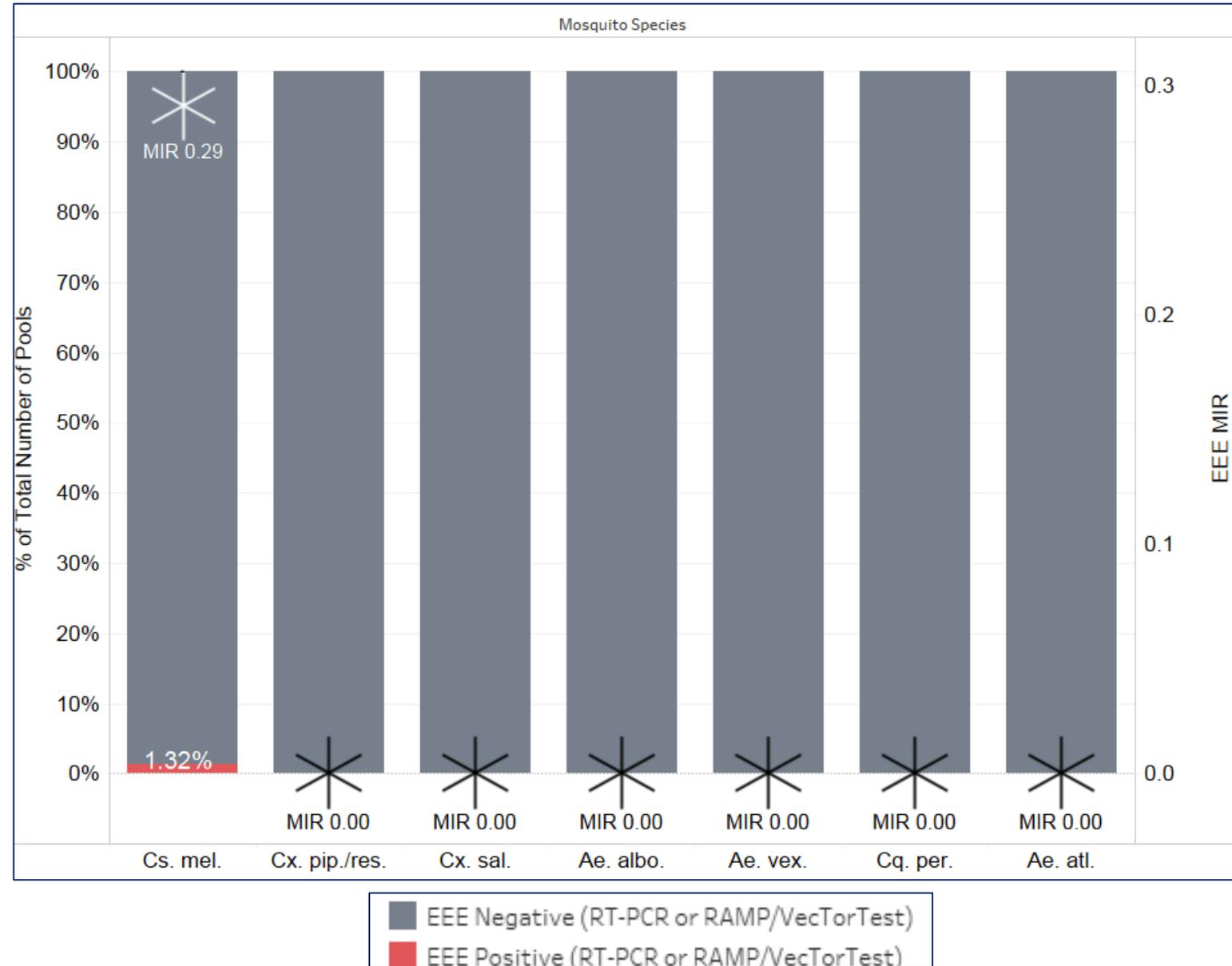
16% of *Cx. pip./res.* pools were WNV positive for a species-specific 4.99 WNV MIR.

Mosquito Species	Number of Mosquitoes Tested	Number of Pools Tested	WNV Positive Pools
<i>Cx. pip./res.</i>	191,173	6,047	953
<i>Ae. albo.</i>	83,372	2,210	21
<i>Cx. spp.</i>	8,892	659	19
<i>Cx. err.</i>	20,051	754	8
<i>Cx. sal.</i>	16,620	641	6
<i>Ae. jap.</i>	628	41	1
<i>Ae. vex.</i>	12,231	484	1
<i>Ae. atl.</i>	97	2	0
<i>Ae. tris.</i>	666	44	0
<i>An. punc.</i>	39	1	0
<i>Cq. per.</i>	378	9	0
<i>Cs. mel.</i>	59,179	1,316	0
Grand Total	393,326	12,208	1,009



1% of *Cs. mel.* pools were EEE positive for a species-specific 0.29 EEE MIR.

Mosquito Species	Number of Mosquitoes Tested	Number of Pools Tested	EEE Binary
<i>Cs. mel.</i>	96,252	2,125	28
<i>Cx. pip./res.</i>	8,517	371	0
<i>Cx. sal.</i>	2,022	46	0
<i>Ae. albo.</i>	989	24	0
<i>Cq. per.</i>	378	9	0
<i>Ae. atl.</i>	97	2	0
<i>Ae. vex.</i>	52	1	0
Grand Total	108,307	2,578	28



- ✓ Complete
- In progress
- !! Goal

Ongoing Projects at VDH

New Arboviral page in statewide disease surveillance system

- Better data collection

Arboviral Investigation Guidelines- Mosquito Control Communication Toolkit

- Provide contact list and protocol for those in jurisdictions with MC
- LHD should notify MCD on positive WNV IgM

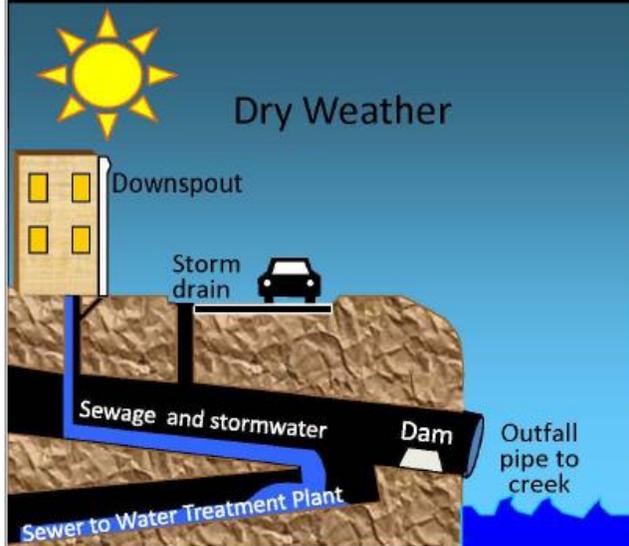
New mosquito surveillance data collection system

- Dropbox sheets

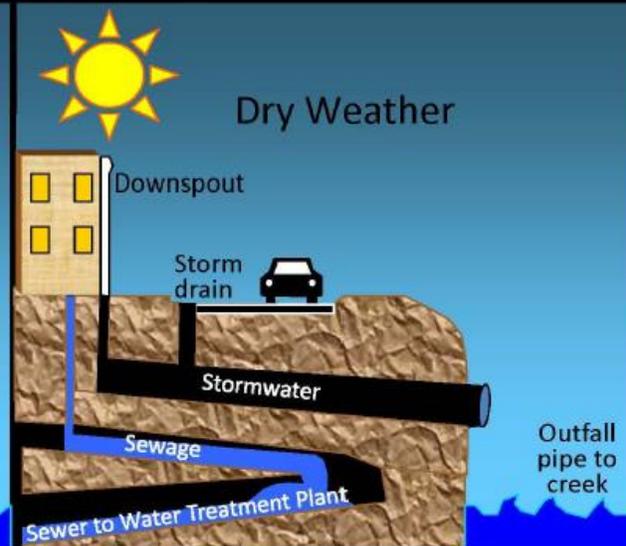
Monthly mosquito surveillance and arboviral disease report on VDH website

- Build public facing interactive and PDF report

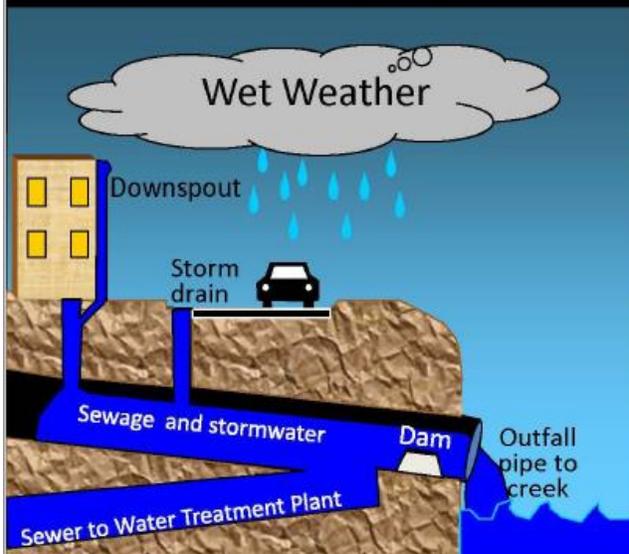
Combined Sewer



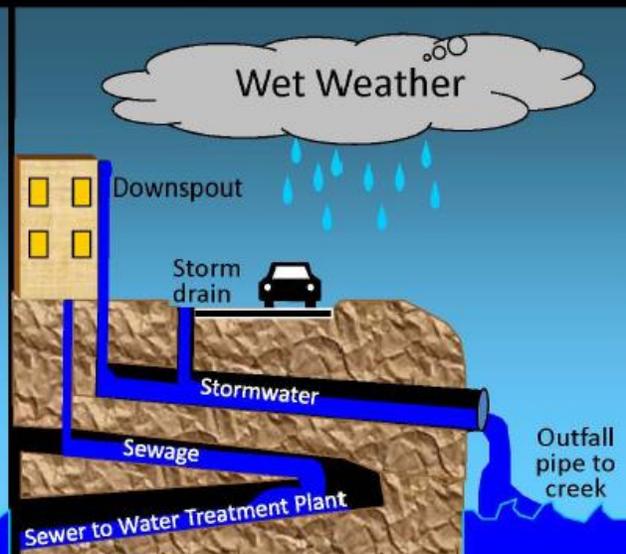
Separate Sewer



Combined Sewer



Separate Sewer



NEVBD Pesticide Resistance Monitoring Program

- The Northeast Regional Center for Excellence in Vector Borne Diseases (NEVBD) is initiating a pesticide resistance monitoring program for the 2019 field season
 - *A specimen submission system will be available, whereby live mosquitoes can be sent directly to Cornell University for resistance testing*
 - *Kits for mosquito collection and pesticide resistance testing will be available*
 - *We will provide education for conducting pesticide resistance testing*
- If you are interesting in participating in our program please visit the NEVBD website: neregionalvectorcenter.com/resistance
 - *The site is still under development, but please include your contact information and we will notify you of program developments*

If you have questions about the program please contact James Burtis

jb766@cornell.edu



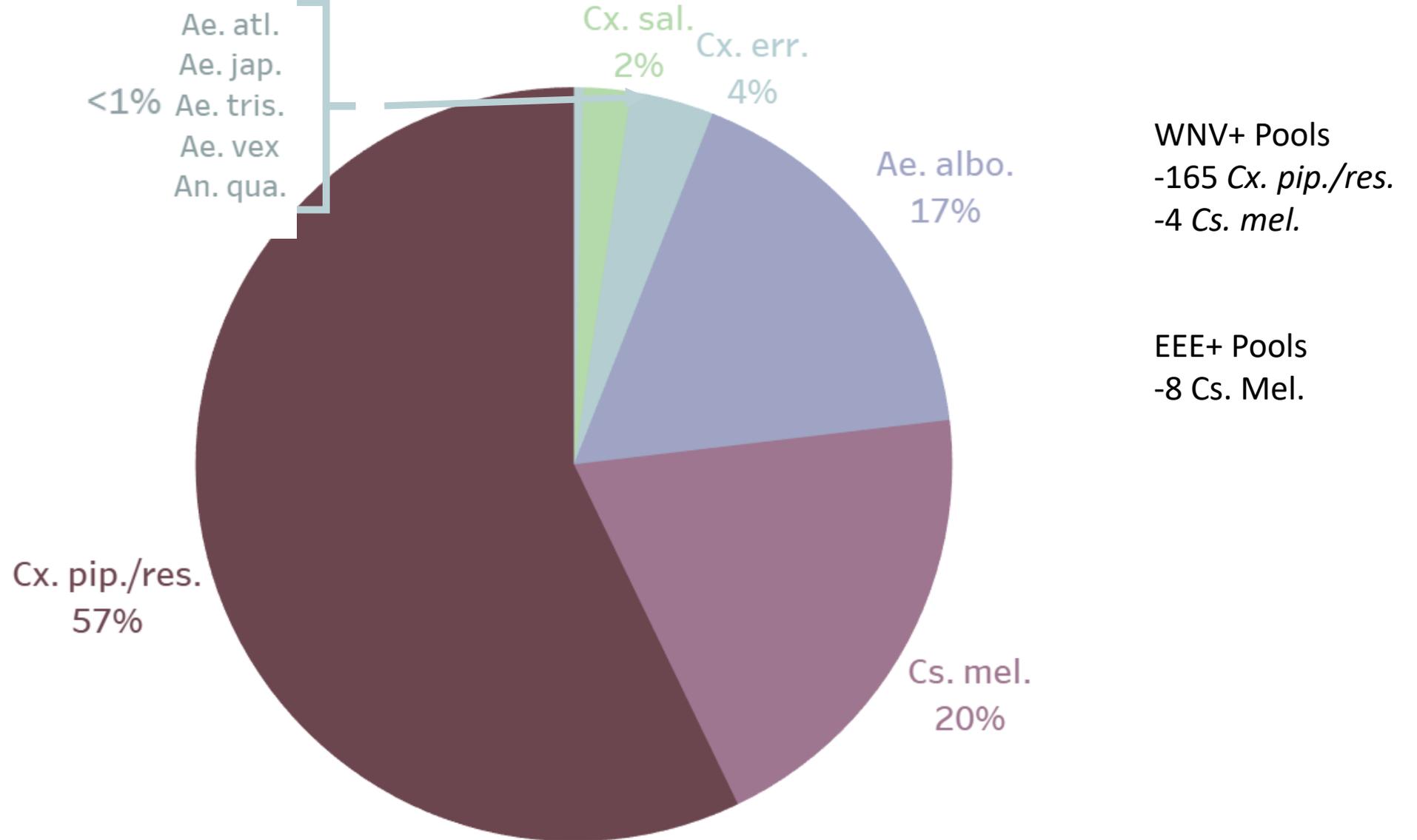
2017

Mosquito Species Tested for WNV	Number of Mosquitoes Tested	Number of Pools Tested	WNV Positive Pools	WNV MIR
<i>Cx. pipiens/restuans</i>	201,641	5,810	744	3.690
<i>Cs. melanura</i>	141,913	2,005	5	0.035
<i>Ae. albopictus</i>	67,905	1,814	8	0.118
<i>Cx. erraticus</i>	21,459	674	3	0.140
<i>Cx. salinarius</i>	13,103	429	8	0.611
<i>Ae. vexans</i>	8,976	323	3	0.334
<i>Cx. spp.</i>	5,477	413	17	3.104
<i>Ae. japonicus</i>	432	29		
<i>Ae. triseriatus</i>	346	25		
<i>An. quadrimaculatus</i>	319	9		
<i>Cq. perturbans</i>	79	2		
<i>An. crucians</i>	41	1		
<i>An. punctipennis</i>	28	1		
Grand Total	461,719	11,535	788	

2018

Mosquito Species	Number of Mosquitoes Tested	Number of Pools Tested	WNV Positive Pools
<i>Cx. pip./res.</i>	191,173	6,047	953
<i>Ae. albo.</i>	83,372	2,210	21
<i>Cs. mel.</i>	59,179	1,316	0
<i>Cx. err.</i>	20,051	754	8
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<i>Ae. atl.</i>	97	2	0
<i>An. punc.</i>	39	1	0
Grand Total	393,326	12,208	1,009

Culex pipiens/restuans, *Aedes albopictus*, & *Culiseta melanura* accounted for 94% of all tested mosquitoes.



Combined Sewage Systems?

