

Developing mosquito-based detection methods for the brown treesnake in Guam



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Mariana Islands













Mariana fruit bat

 Endemic to Mariana and Caroline Islands

Federally endangered





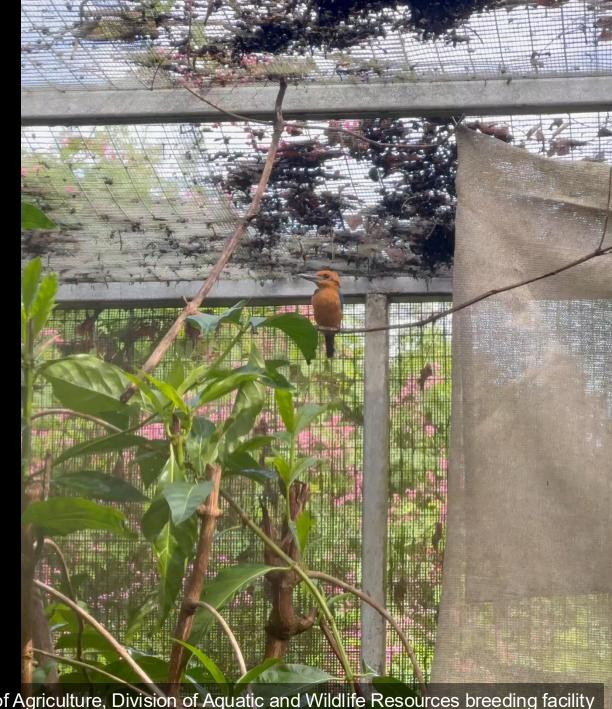


Guam kingfisher

- **Endemic to Guam**
- Extinct in the wild since 1986
- Global population: <200



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Bridled white-eyeGuam subspecies extinct
Last seen 1983



Mariana crow
Critically endangered
Extirpated from Guam



Guam flycatcher Extinct Last seen 1983



Micronesian rufous fantail Guam subspecies extinct Last seen 1984



Mariana fruit dove Extirpated from Guam Last seen 1984



Micronesian honeyeater
Extirpated from Guam
Last seen 1986

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Brown treesnake Boiga irregularis





- Mildly venomous colubrid from northern Australia and Papua New Guinea
- Introduced to Guam shortly after WWII, likely from Papua New Guinea
- First detection in the early 1950s, island-wide by 1968
- No natural predators, abundant, naïve prey species
- Extreme densities: >13,000 snakes per sq. mile, exceeding carrying capacity
- Extirpation of most of Guam's forest vertebrates, extinction of 12 native birds
- Widespread power outages; ~150 per year due to snakes
- Brown tree snake populations currently decreasing on Guam





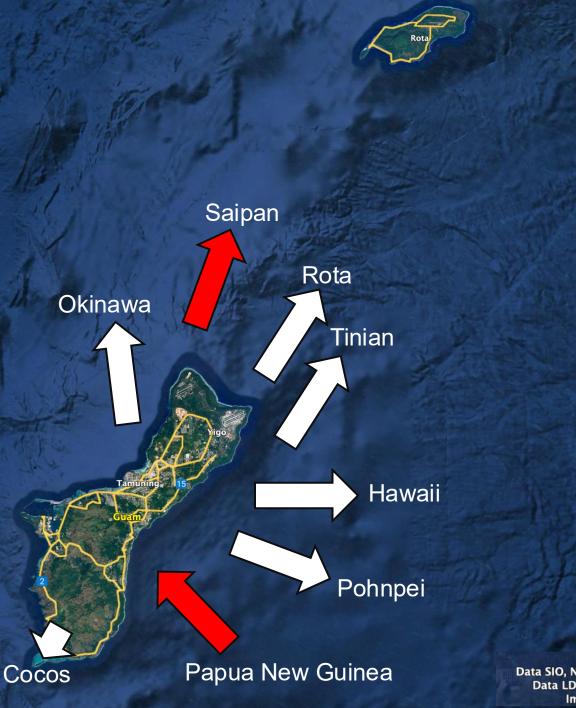




Dispersal of snakes from Guam

| Destination | Number intercepted (2009) |
|--------------------------------|------------------------------|
| Rota | 4 |
| Saipan | 78 |
| Tinian | 8 |
| Diego Garcia | 1 |
| Federated States of Micronesia | 1 |
| Palau | 1 |
| United States Mainland | 3 |
| Hawaii | 14 |

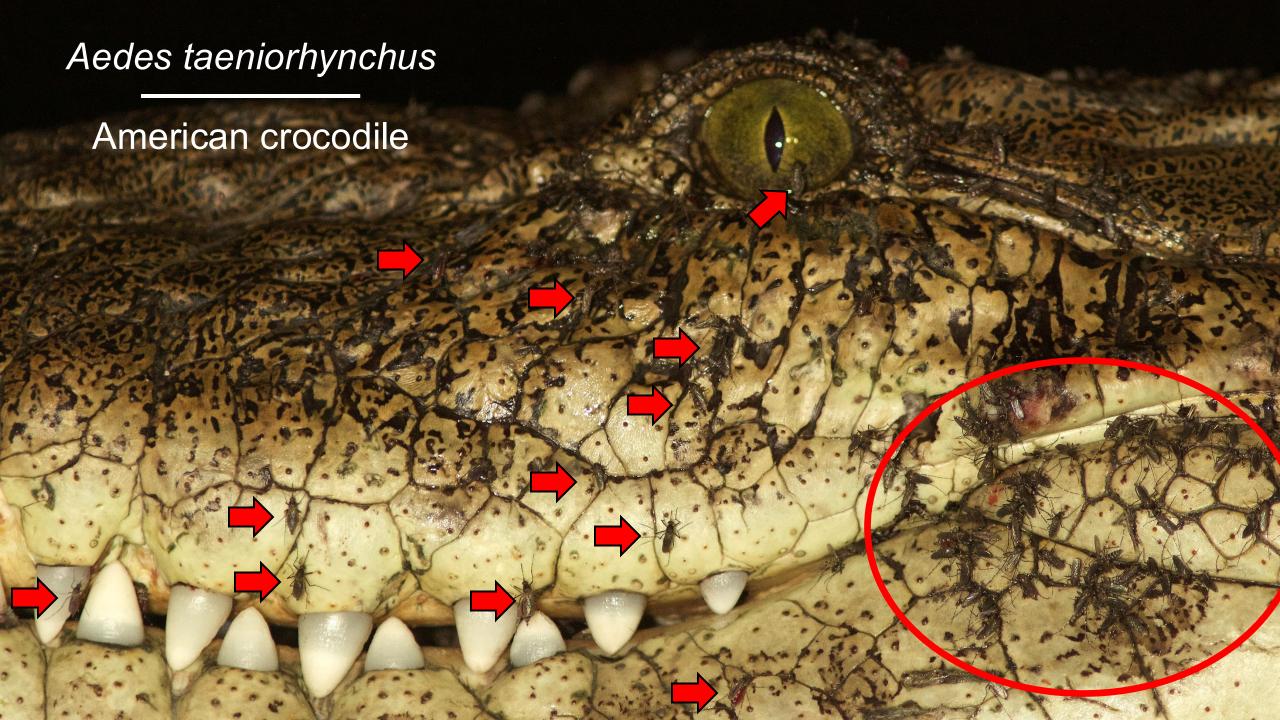
Perry and Vice 2009



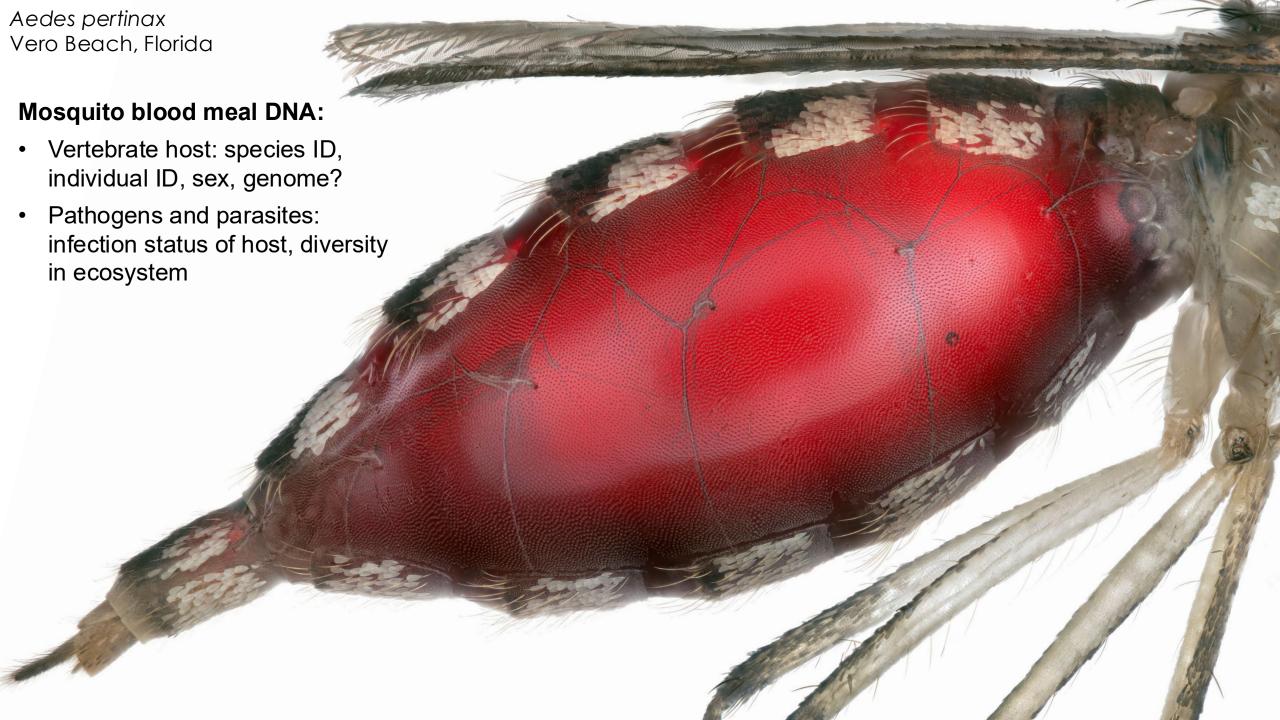
Data NOAA Data SIO, NOAA, U.S. Navy, NGA, GEBCO Data LDEO-Columbia, NSF, NOAA Image © 2025 Airbus How can we detect incipient populations of brown treesnakes in new locations?

How can we detect incipient populations of brown treesnakes in new locations?

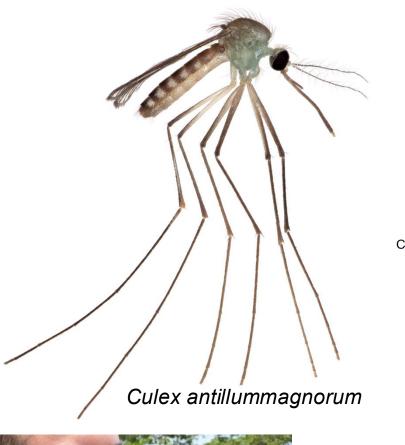




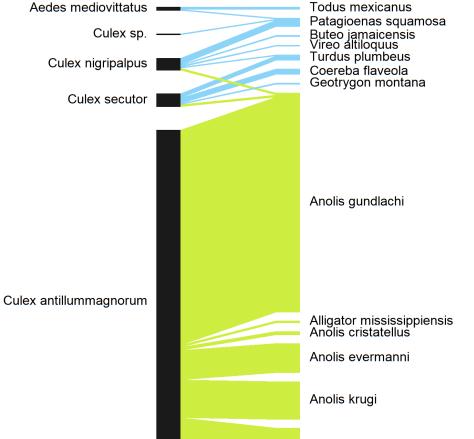


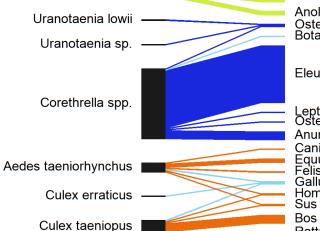


Plasmodium diversity

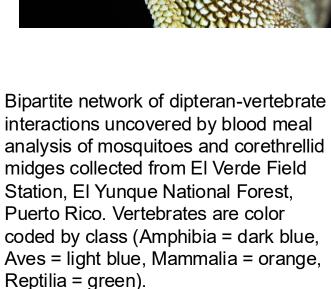










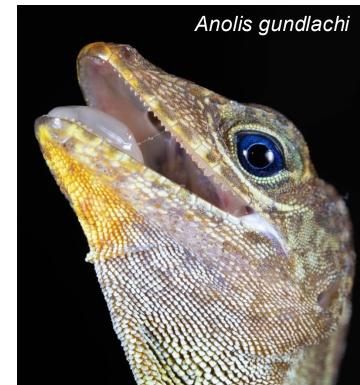


Anolis cuvieri Osteopilus septentrionalis Botaurus lentiginosus

Anolis stratulus

Eleutherodactylus spp.

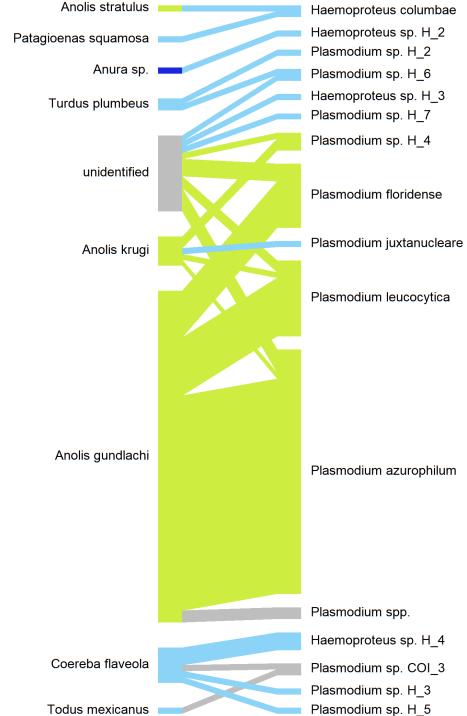
Leptodactylus fuscus Osteocephalus helenae Anura spp. Canis familiaris Equus caballus Felis catus Gallus gallus Homo sapiens Sus scrofa Bos taurus Rattus norvegicus

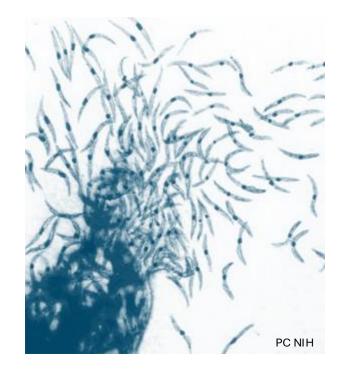


Plasmodium diversity









Plasmodium azurophilum

Plasmodium spp.

Haemoproteus sp. H 4

Plasmodium sp. COI 3

Plasmodium sp. H 3 Plasmodium sp. H 5

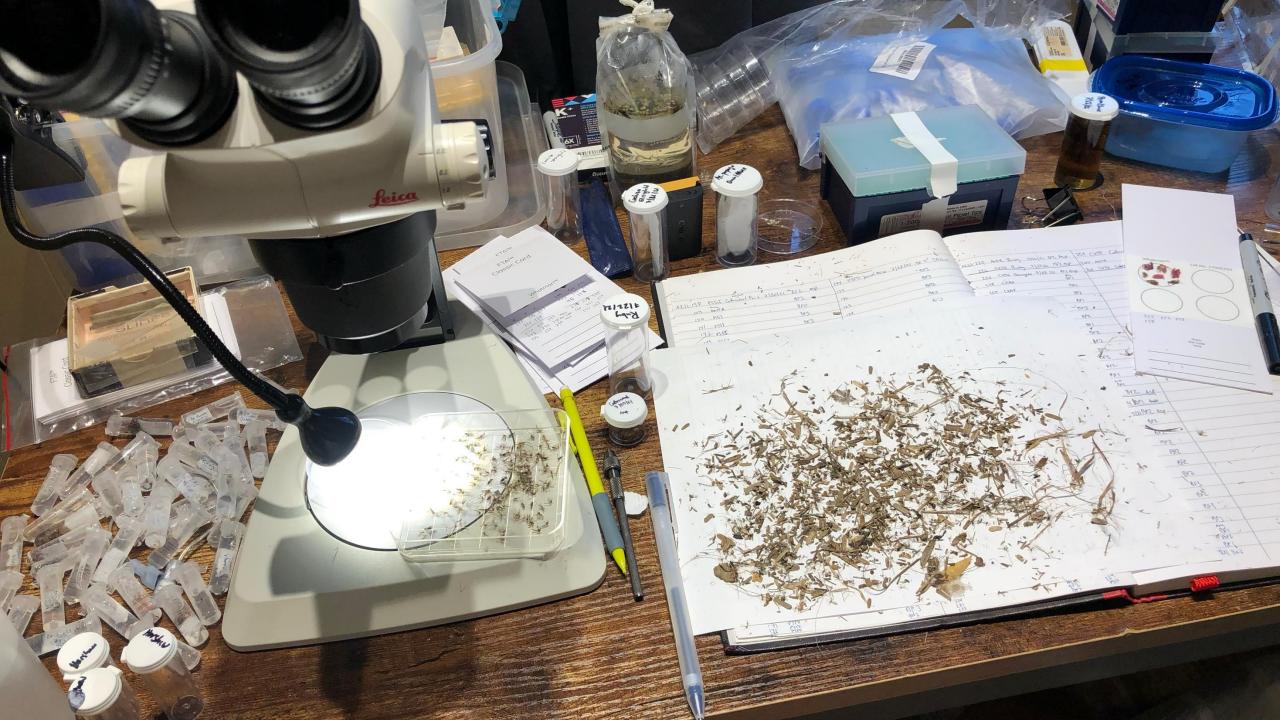
Bipartite network of vertebrate-parasite interactions uncovered by haemosporidian-targeting blood meal analysis of mosquitoes and corethrellid midges collected from El Verde Field Station, El Yunque National Forest, Puerto Rico. Vertebrates are color coded by class and parasite species/OTUs are color coded by class of host (Amphibia = dark blue, Aves = light blue, Mammalia = orange, Reptilia = green, unknown = grey).

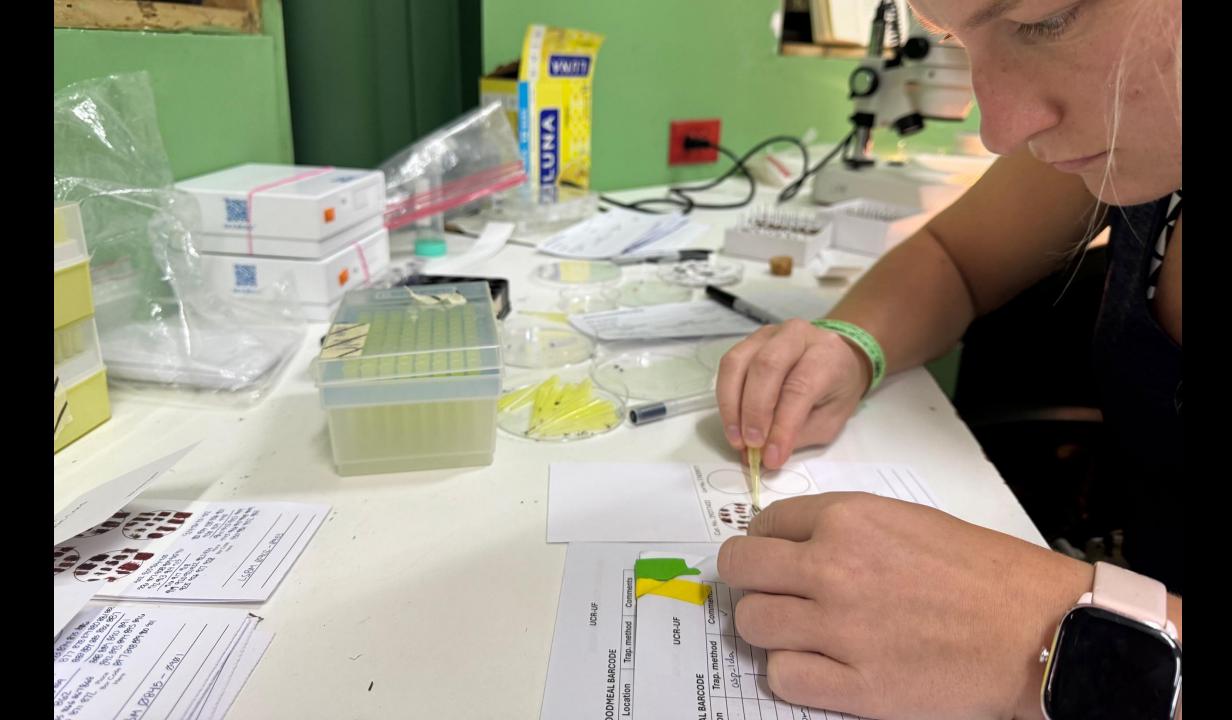


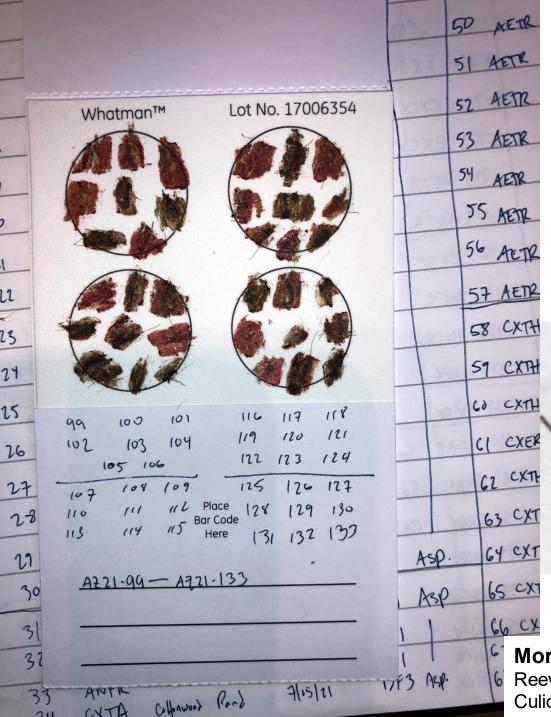












Field preservation of mosquito blood meals on Whatman FTA Classic Cards

- Effective DNA preservation for extended duration
- Room temperature storage
- No specialized equipment necessary
- RNA expected to be preserved as well



More information:

Reeves et al. 2016. Maintenance of host DNA integrity in field-preserved mosquito (Diptera: Culicidae) blood meals for identification by DNA barcoding. Parasites and Vectors 9: 503.

Blood Meal Analysis:

A collection of molecular techniques designed to taxonomically identify the source of an invertebrate's blood meal

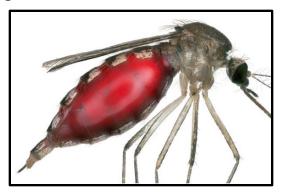
Serological assays (ELISA, precipitin test)

DNA-based: PCR, PCR + Sanger sequencing

Some next gen sequencing approaches

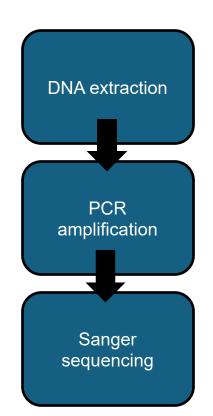
DNA barcoding workflow

Specimen collection



- Fresh specimens
- Preserved specimens
- Museum specimens
- Mosquito blood meals

Laboratory processing



Sequence Identification





Deluca 3366-3409

mosquito blood meals. PLOS Negl. Trop. Dis. 12: e0006767.

DELU3158 - 3/96

DeLva 3454-3497

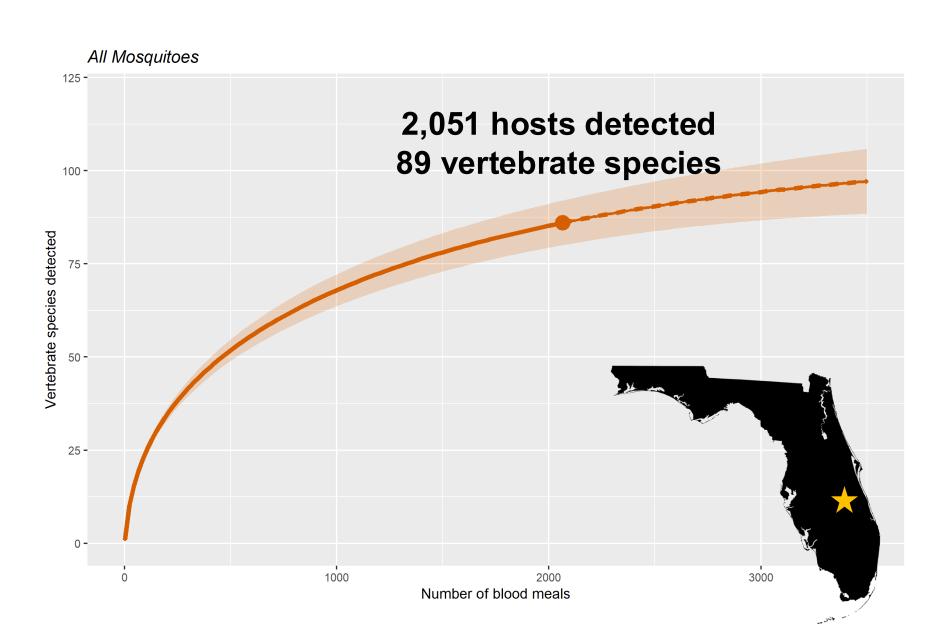
Deluca 3410 - 3453

Mosquito blood meal-based vertebrate detection, DeLuca Preserve, Osceola Co., Florida

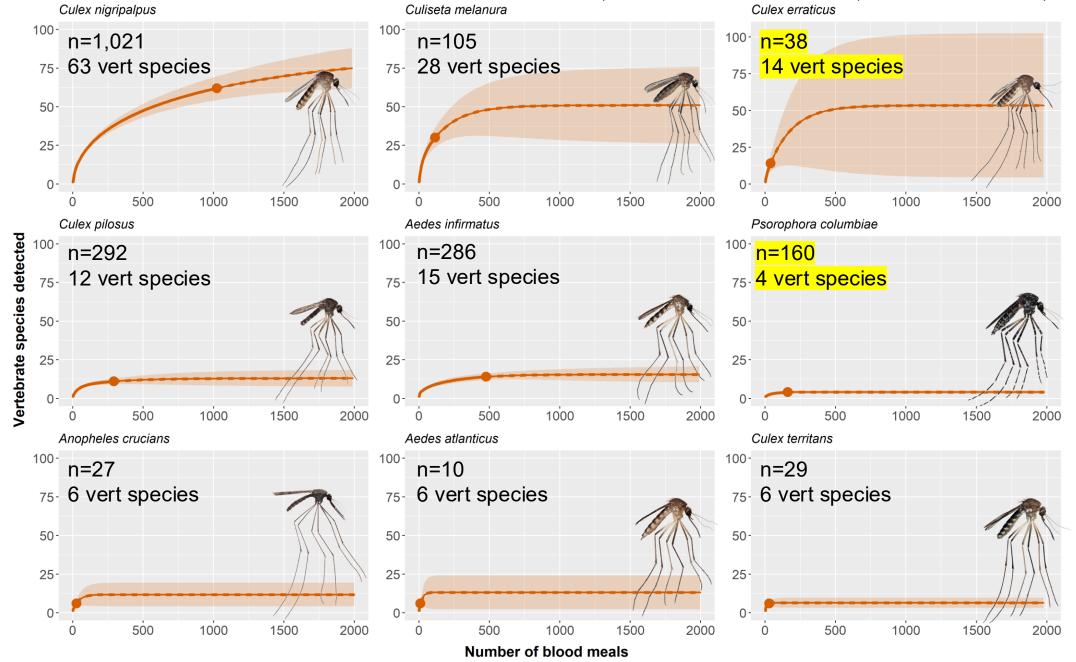


Hannah Atsma

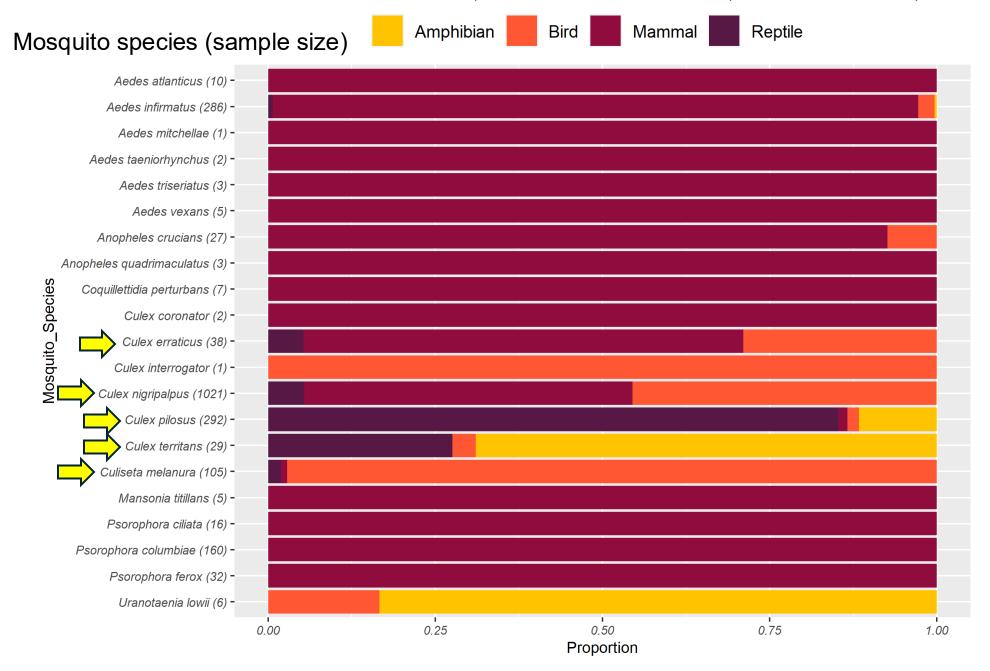
- Mosquitoes collected across nine months
- 54,637 mosquitoes collected
- 3,508 blood meals
- 21 mosquito species



Mosquito blood meal-based vertebrate detection, DeLuca Preserve, Osceola Co., Florida



Mosquito blood meal-based vertebrate detection, DeLuca Preserve, Osceola Co., Florida



5/21 mosquito species fed on reptilian hosts

Goal: Develop mosquito blood meal-based methods for detecting brown tree snakes in Guam and the Marianas



- 1. Collect blood fed mosquitoes and identify species likely to feed on reptiles in the Marianas
- 2. Develop resources for mosquito identification and recognition of reptile-feeding species
- 3. Train local personnel on methods of collecting blood fed mosquitoes and processing mosquito blood meals

Mosquito fauna of Guam

Native, endemic species

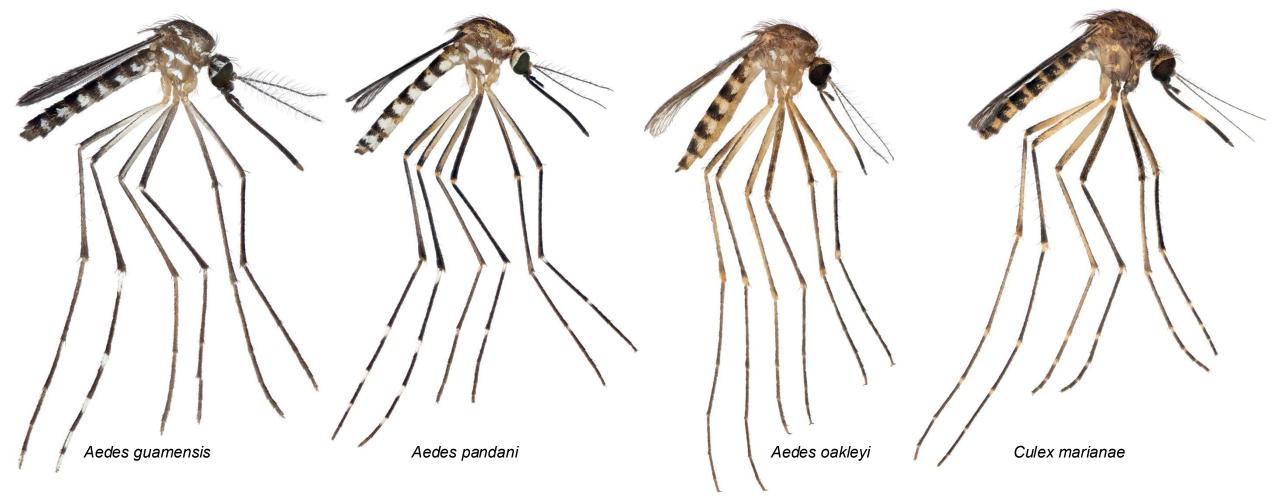
1. Aedes (Stegomyia) guamensis

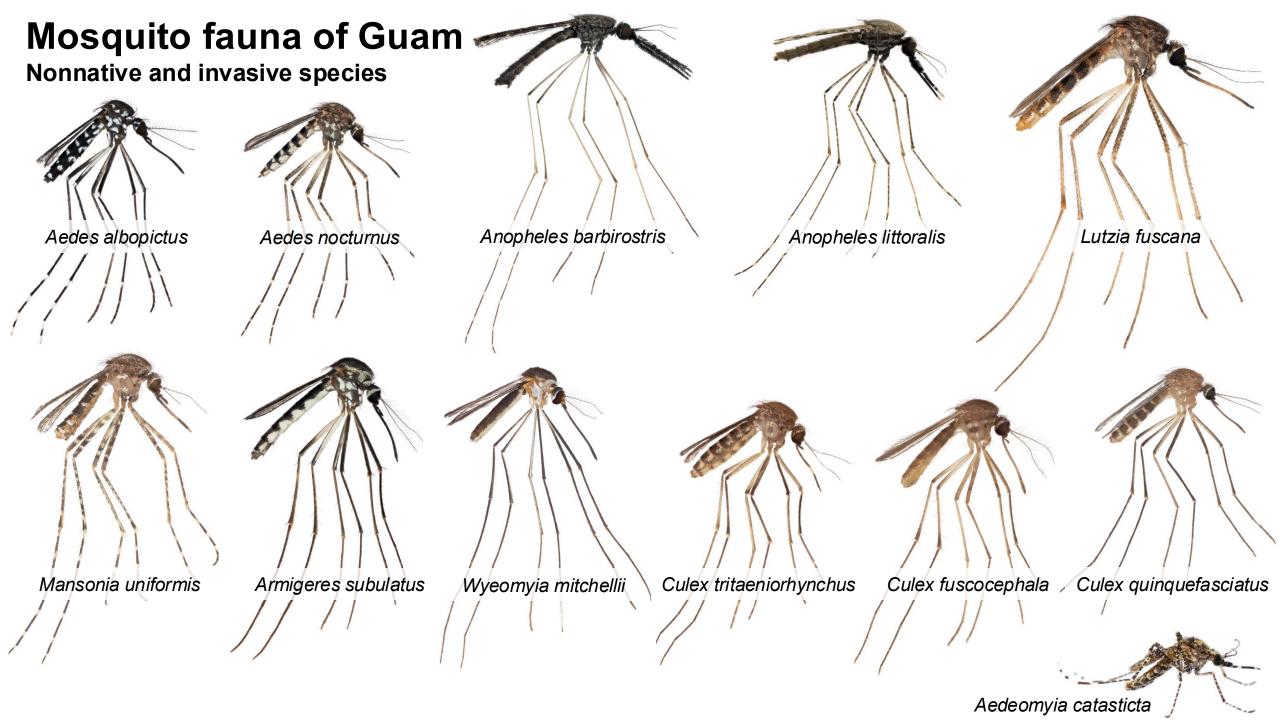
Pandani Group

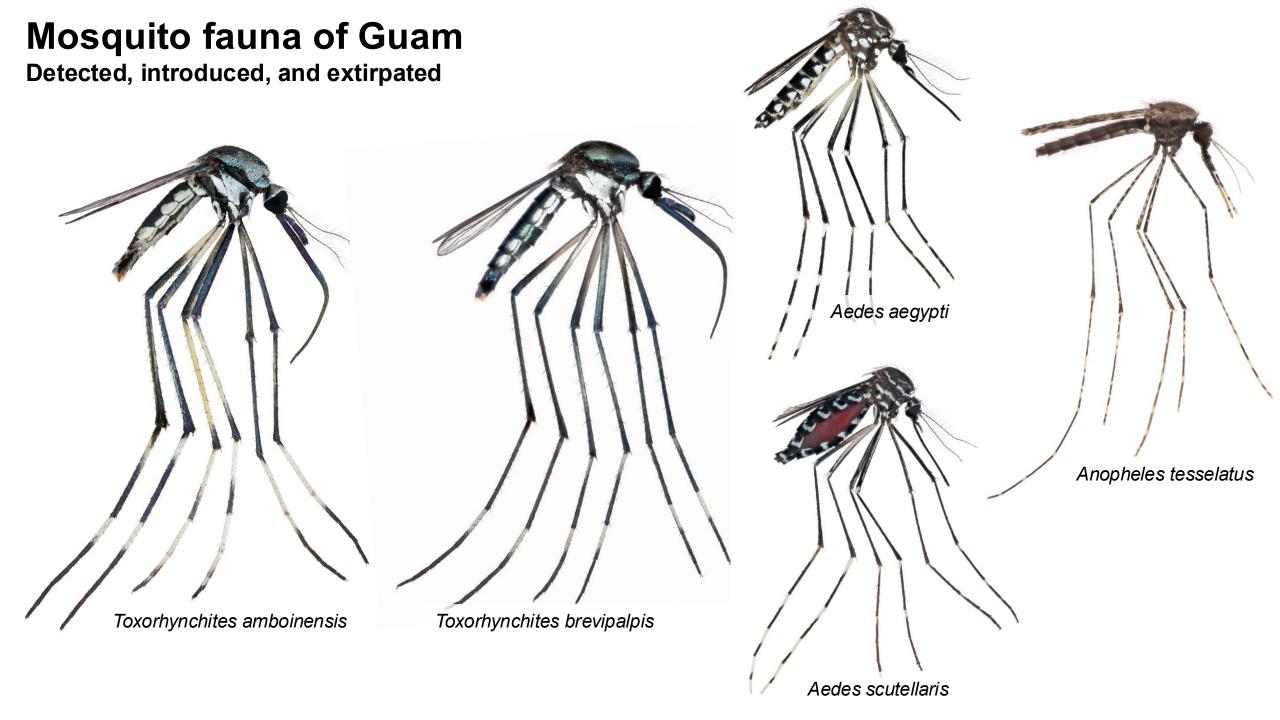
- 2. Aedes (Stegomyia) pandani
- 3. Aedes (Stegomyia) rotanus
- 4. Aedes (Stegomyia) saipanensis
- 5. Aedes (Aedimorphus) oakleyi

Genus Culex

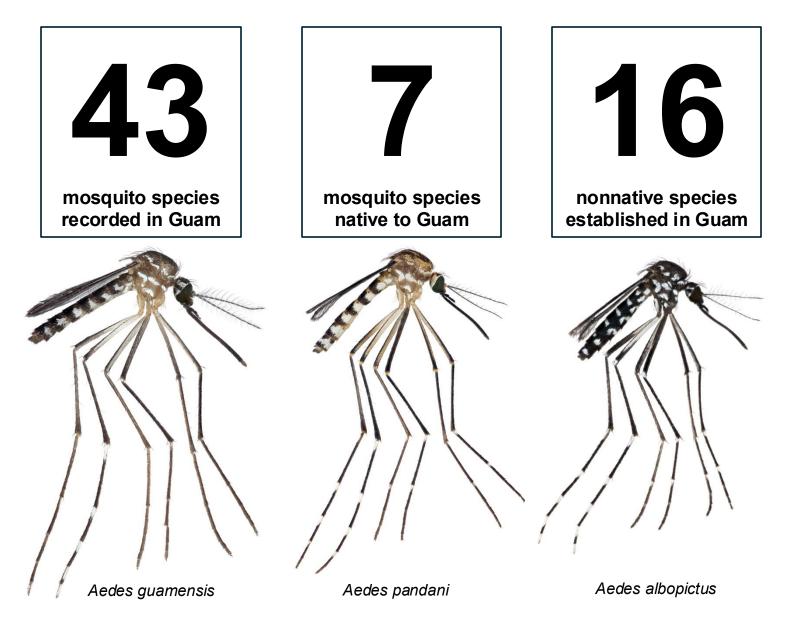
- 6. Culex (Culex) marianae
- 7. Culex (Culex) litoralis







Mosquito fauna of Guam



Host associations

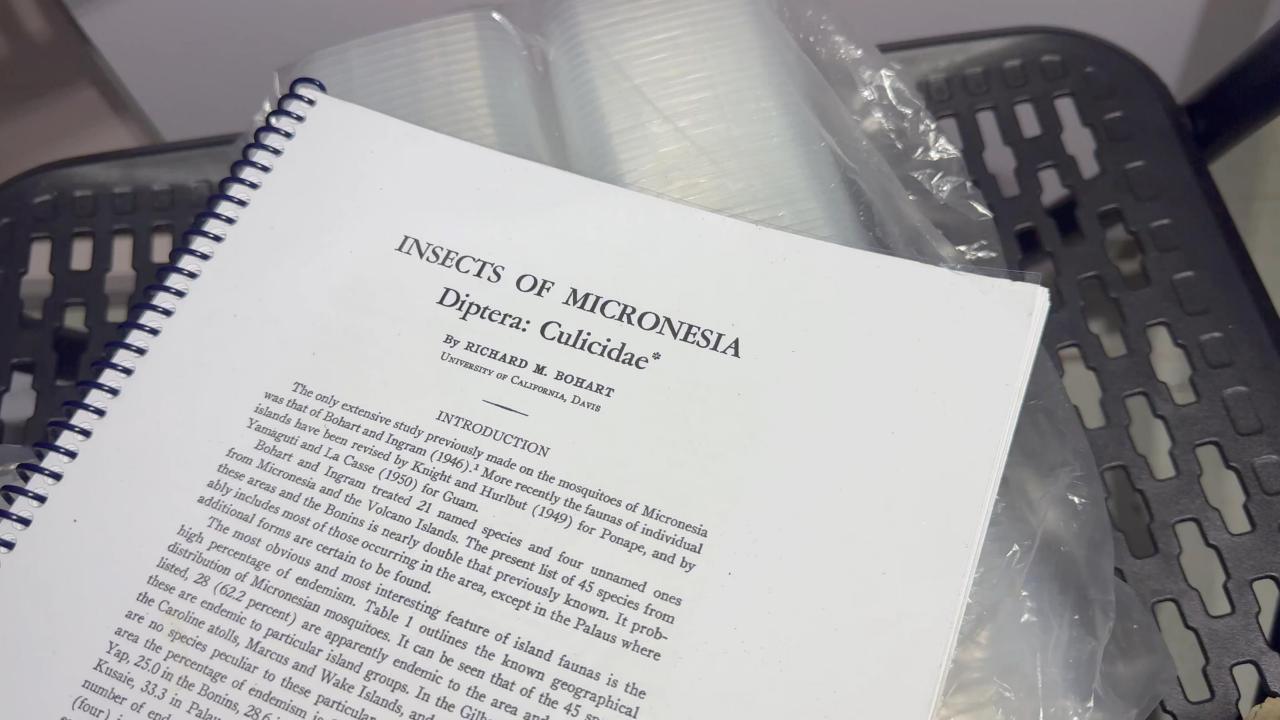
- Mosquito host associations have never been assessed in wild mosquitoes in Guam
- Reeves and Rudnick 1951
 assessed human feeding with
 indoor collections of four species
- Host associations of native species, others, unknown

Mosquito collections, Oct. 2024 Guam Cocos

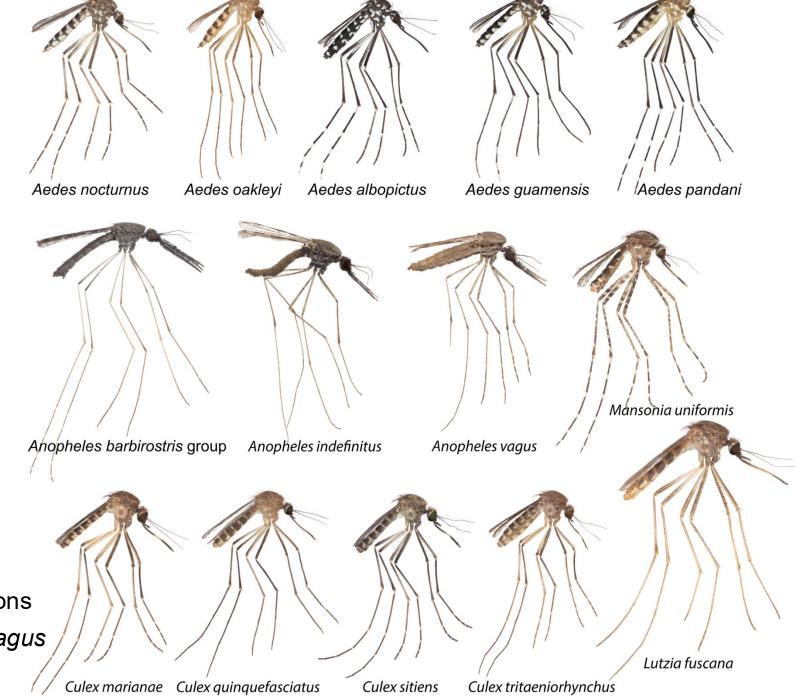






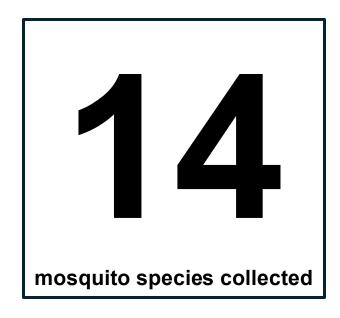


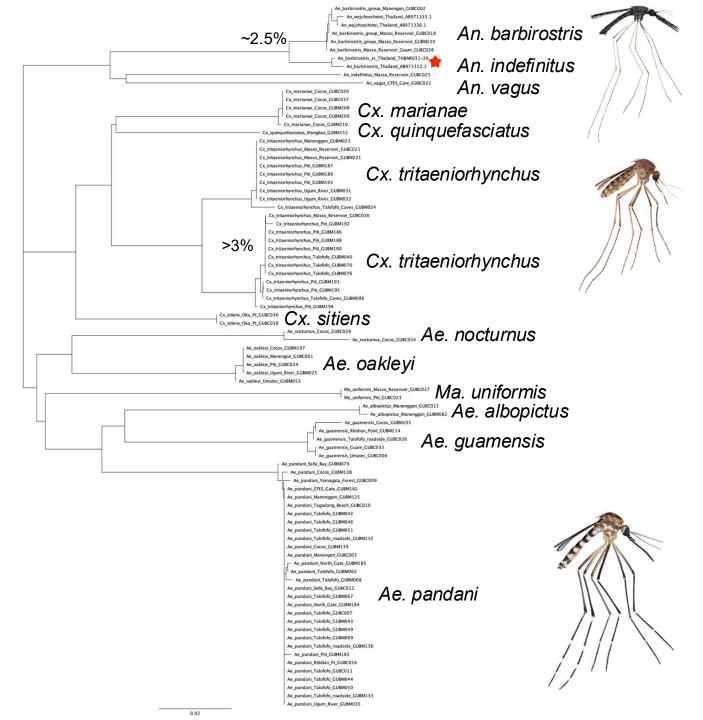
mosquito species collected



*Culex sitiens only found in larval collections

**Anopheles indefinitus and Anopheles vagus only collected in CDC light traps





^{*}Culex sitiens only found in larval collections

**Anopheles indefinitus and Anopheles vagus
only collected in CDC light traps

Mosquito blood meals collected:

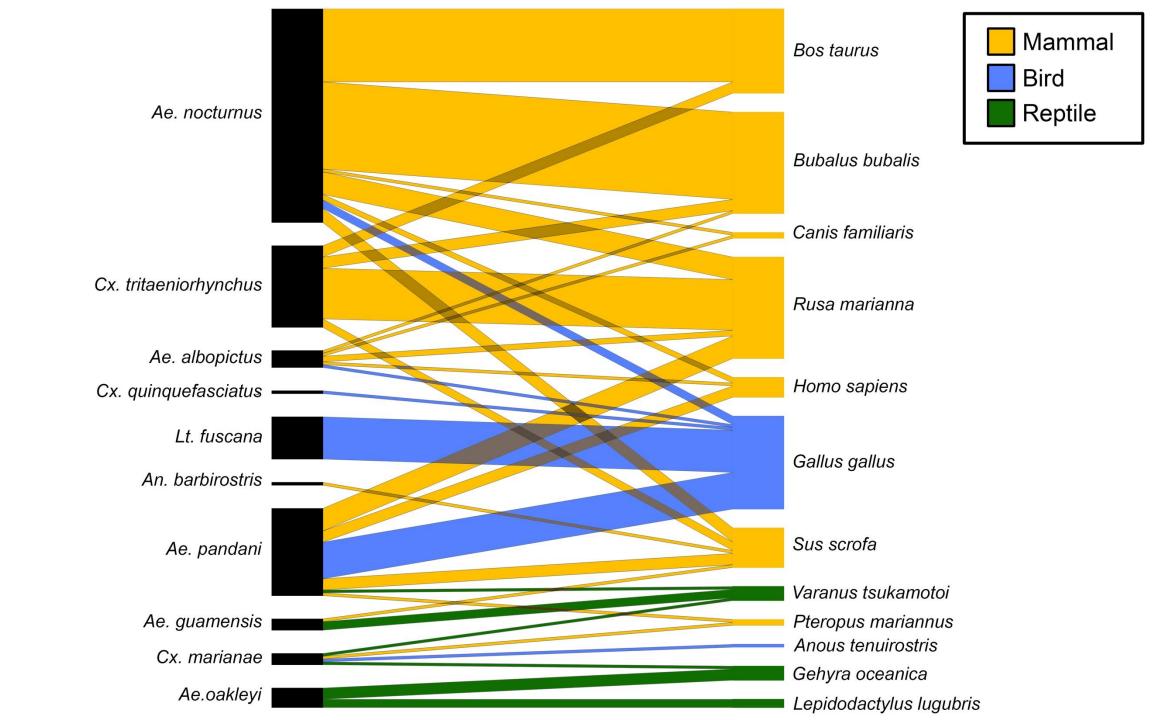
221

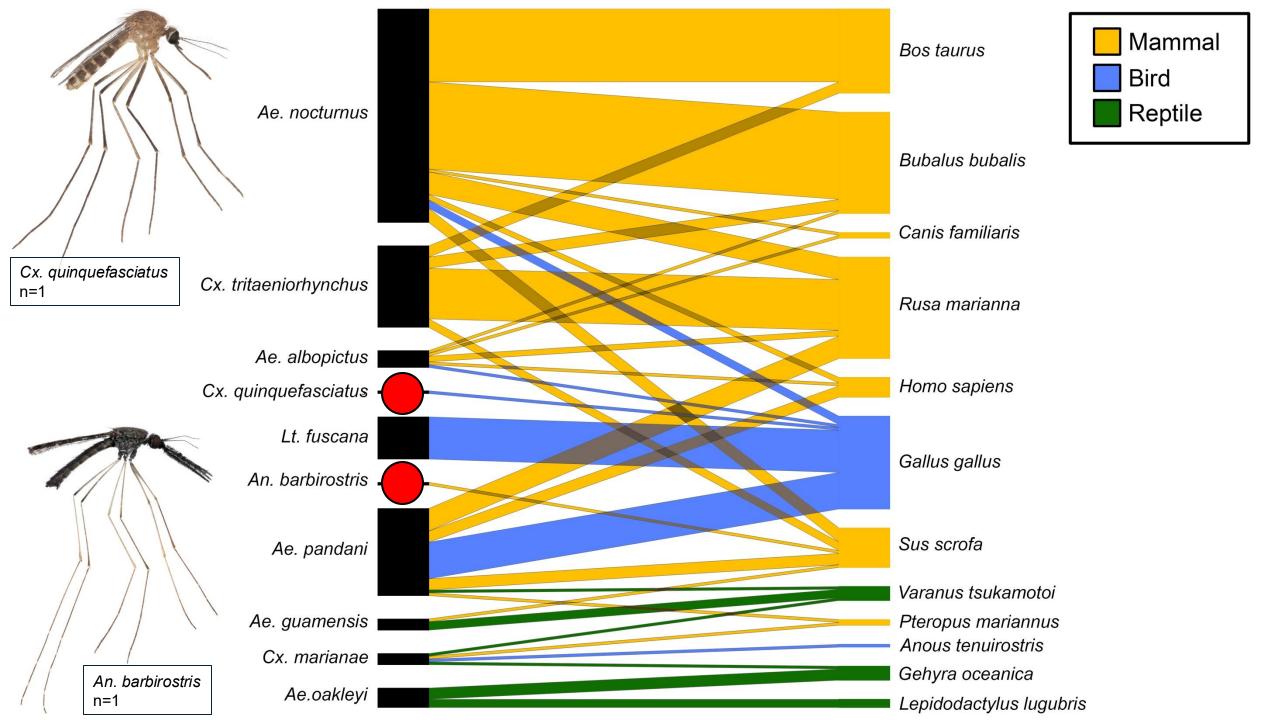
Mosquito blood meals identified:

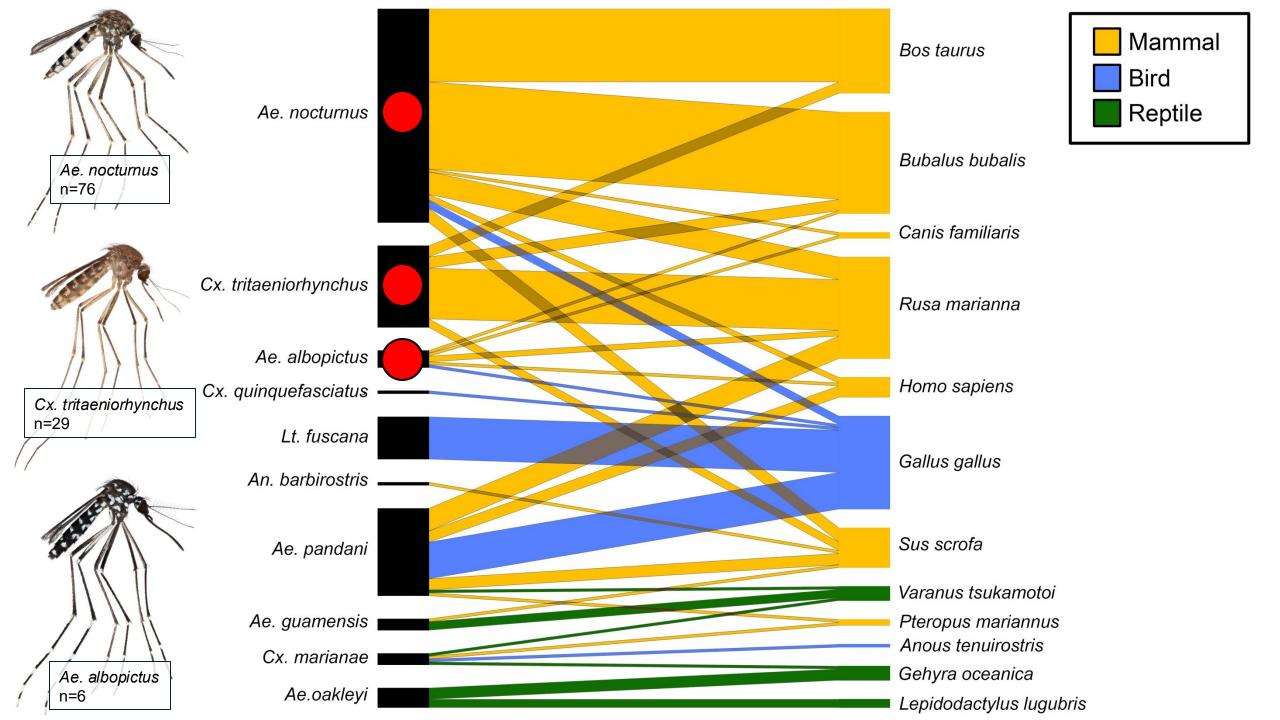
174 (79%)

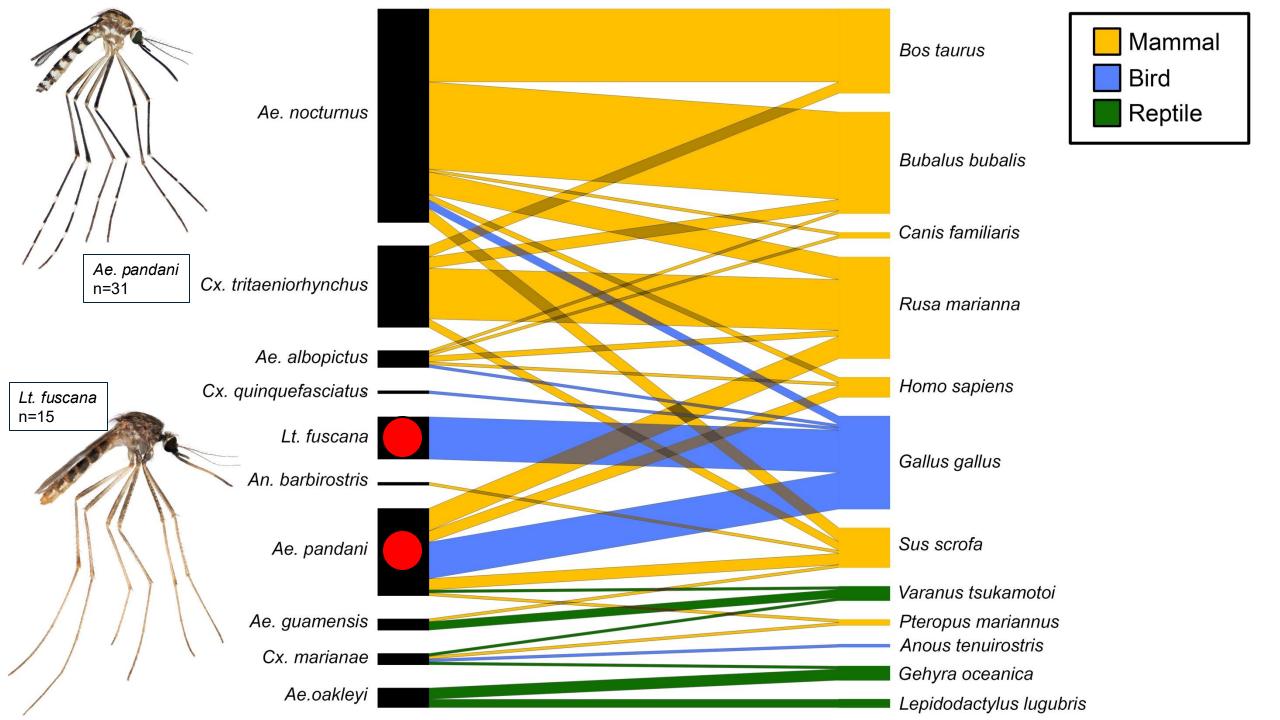


| Mosquito species | Number of blood meals | Number identified (%) |
|------------------------------------|-----------------------------|-----------------------------|
| Aedes (Stegomyia) albopictus | 7 | 6 (86%) |
| Aedes (Stegomyia) guamensis | 5 | 4 (80%) |
| Aedes (Stegomyia) Pandani Group | 42 | 31 (74%) |
| Aedes (Aedimorphus) nocturnus | 83 | 76 (92%) |
| Aedes (Aedimorphus) oakleyi | 14 | 7 (50%) |
| Anopheles (Anopheles) barbirostris | 1 | 1 (100%) |
| Culex (Culex) quinquefasciatus | 2 | 1 (50%) |
| Culex (Culex) tritaeniorhynchus | 38 | 29 (76%) |
| Culex (Culex) marianae | 6 | 4 (67%) |
| Lutzia (Metalutzia) fuscana | 23 | 15 (65%) |

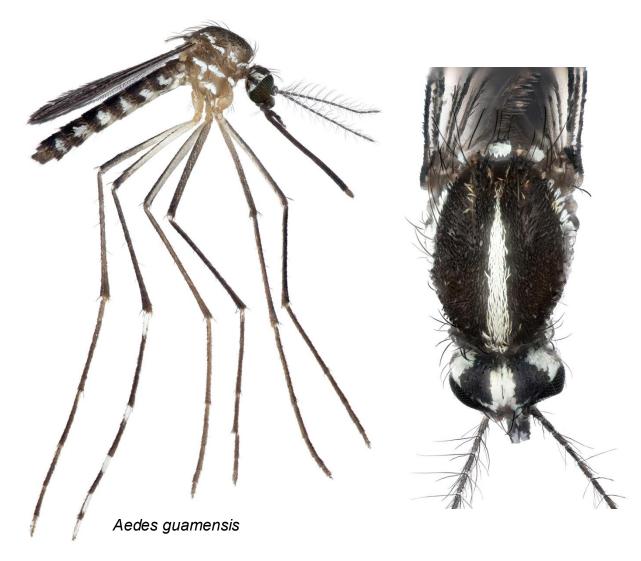








Aedes (Stegomyia) guamensis

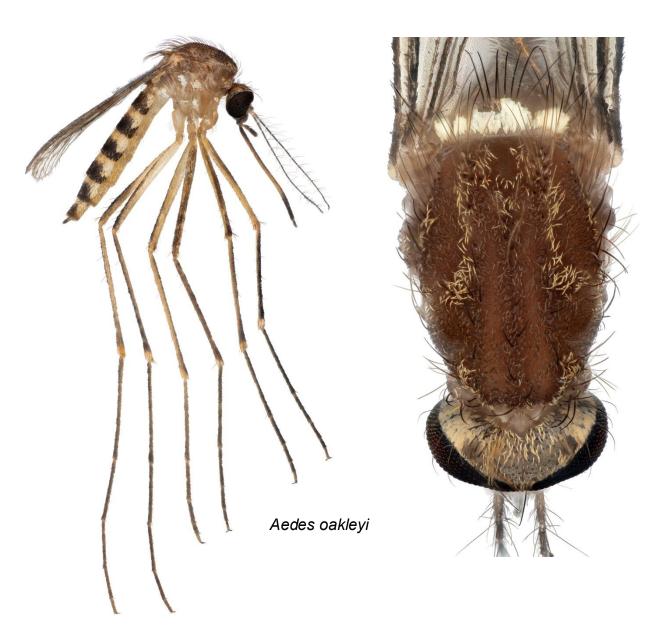


- Member of subgenus *Stegomyia*, like *Ae. aegypti* and *Ae. albopictus*
- Native, endemic to Mariana Archipelago
- Dominant tree hole mosquito prior to arrival of Ae. albopictus
- Larvae in natural and artificial containers
- Host associations previously unknown, does not bite humans
- Strong association with reptiles (3/4 bood meals from monitor lizards)



Larval habitat of Aedes guamensis, Cocos

Aedes (Aedimorphus) oakleyi



- Member of subgenus Aedimorphus, like Ae. vexans
- Native, endemic to Mariana Archipelago
- Host associations previously unknown, does not bite humans
- Larvae in natural and artificial containers
- Poorly known, resting sites unstudied
- Strong association with reptilian hosts (7/7 blood meals from geckos), likely a specialist of reptiles



Culex (Culex) marianae



- Member of subgenus *Culex*, like *Cx*. quinquefasciatus
- Previously a subspecies of *Cx. annulirostris*
- Native, endemic to Mariana Archipelago
- Host associations previously unknown, not strongly attracted to human hosts
- Larvae in fresh to brackish ground pools, containers, vegetated water
- Small sample size, but appears to be a generalist









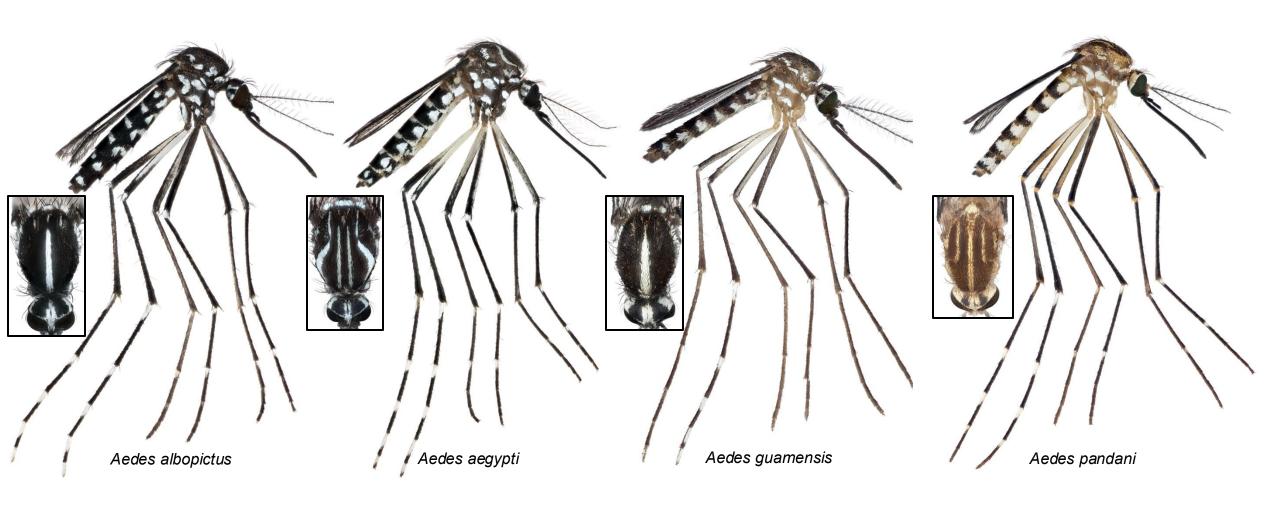
Lesser noddy

Oceanic gecko

Mariana monitor

Mariana fruit bat

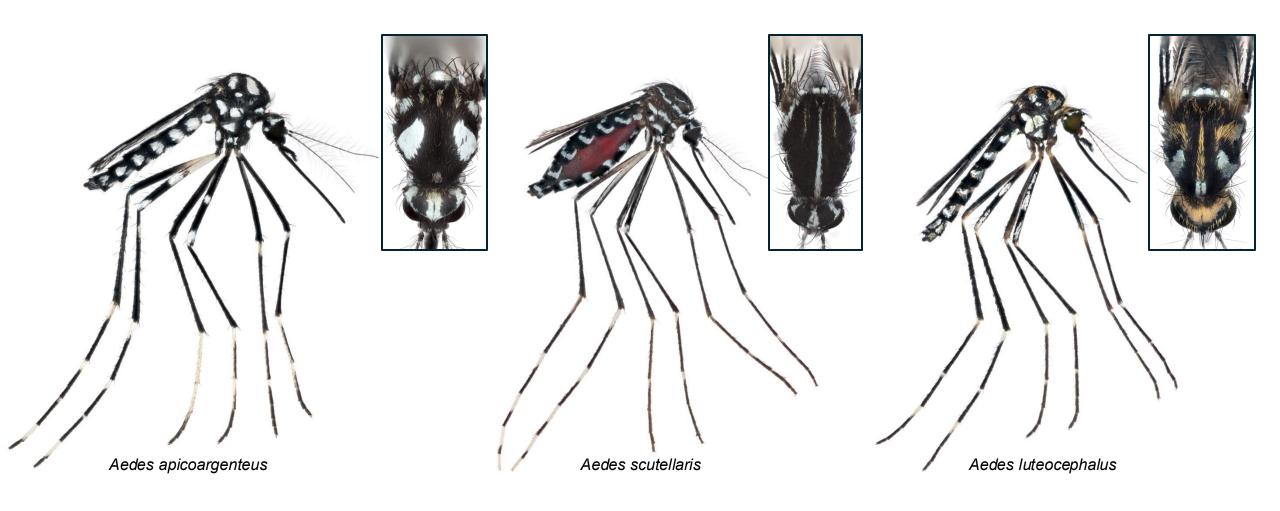
Aedes subgenus Stegomyia



Widespread/cosmopolitan, globally important vectors

Poorly known, Mariana Archipelago endemics

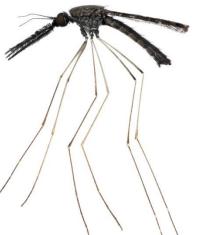
Aedes subgenus Stegomyia – 133 species, globally



Conclusions

- Aedes guamensis and Ae. oakleyi are likely specialists of reptilian hosts, but sample size was limited
- Culex marianae is likely a generalist that will feed from reptiles, but sample size was limited
- Aedes nocturnus, Ae. pandani, and Cx. tritaeniorhynchus are abundant, but feed from mammals or birds (Ae. pandani)
- Aedes guamensis, Aedes oakleyi, Culex marianae are candidate species for maximizing return on investment in future surveys for brown treesnakes on Guam
- Cryptic diversity in Culex tritaeniorhynchus in Guam
- COI sequences from Guam Anopheles barbirostris do not match An. barbirostris s.s.; likely An. wejchoochotei





Acknowledgements

- Julie Siegel, NWRC Grants Specialist, and NWRC leadership for expediting the cooperative agreement
- Dr. Aubrey Moore, University of Guam (retired) for project inspiration
- Alex Urlaub, Jen Espineira, Ray Gellner for molecular lab support
- This work was supported by funding provided by the Department of the Navy, Joint Region Marianas, under IAA# 19-1740-1063-IA. Project management provided by NAVFAC Marianas









