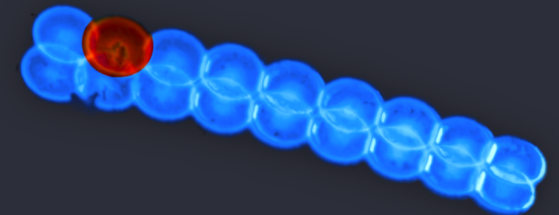
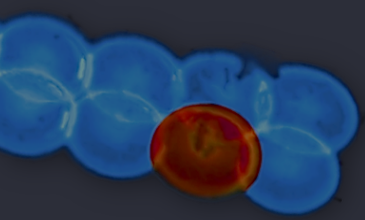


Project PREMONITION

CPS for Disease Surveillance

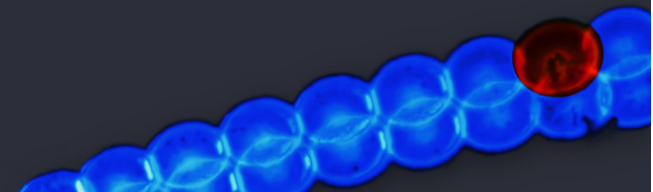
Ethan Jackson – Microsoft Research





Emerging Infectious Diseases

Technologies to stop them.



The Threat of Disease Emergence

Emerging infectious diseases pose significant health and economic threats.

**SARS in China
(2003):**

\$6.2 billion / **8000** = **\$775,000**
cost cases

H5N1 (2003-2009):

\$20 billion / **468** = **\$43 million**
cost cases

**Ebola in US
(2014):**

\$2.8 billion / **4** = **\$700 million**
cost cases

Need Better Surveillance of Diseases

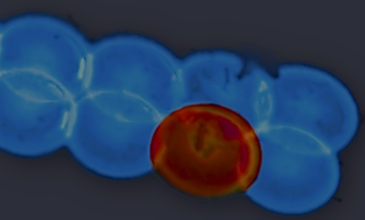
But surveillance of EIDs is particularly challenging because:

- Majority of threats caused by animal pathogens
Over 60% of emergence events
- Best existing systems detect outbreaks too late
Median delay of 13.5 days
- Many emerging diseases were previously unknown
Over 70% of viruses in wild are unknown

Actually Need Surveillance in The

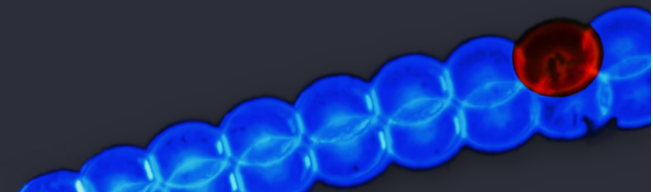
To see which pathogens are circulating in animals before they impact humans

Spoiler: Put us on a path towards an ambitious, massive, self-adapting CPS



Project PREMONITION

Preventative Monitoring of Infectious Agents



Mosquito-as-a-Device

Use mosquitoes as devices that collect blood samples from animals in the wild



Mosquitoes are used as devices that collect blood samples from animals in the wild. This technology has been used in various parts of the world, including Antarctica.

Classical genetic markers are used to identify individual animals, and pathogen analysis is used to identify diseases. This information is used to study the impact of climate change on wildlife and plants. This is done on a large scale.

Mosquito-as-a-Device

Use mosquitoes as devices that collect blood samples from animals in the

- Mosquitoes grow naturally in ^{wild} rural and urban environments
On average live 20 days, consume 2.5 μ l per blood meal, can fly several miles, and are geographically widely distributed. They have advanced olfactory systems to locate hidden prey.
- Mosquitoes sample the genes of animals and their pathogens
In studies, over 70% of viral pathogens in mosquitoes are unknown to science and come from a wide range of hosts, including humans, ducks, geese, cows, and plants.
- Can leverage classical entomological methods
Classical field entomology has been used by researchers, governments, and militaries to efficaciously surveil pathogens carried by mosquitoes. Must re-invent these methods to achieve scaled required for PREMONITION.



Key Innovations

A number of technological innovations are required for scalability

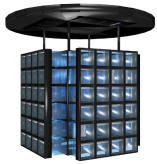


Should be able to read genomic data directly to the computer without the need for a separate storage device

Existing technologies for sequencing DNA are slow and expensive. To read a single human genome, it takes about 60 days and costs about \$100 million. To read a mouse genome, it takes about 10 days and costs about \$10 million. To read a bacterial genome, it takes about 1 day and costs about \$1 million. To read a viral genome, it takes about 1 hour and costs about \$100,000. To read a single human genome, it takes about 60 days and costs about \$100 million. To read a mouse genome, it takes about 10 days and costs about \$10 million. To read a bacterial genome, it takes about 1 day and costs about \$1 million. To read a viral genome, it takes about 1 hour and costs about \$100,000. require manual processing.

Key Innovations

A number of technological innovations are required for scalability



- Smarter Traps that automate field biology

Existing mosquito traps are too low-throughput. They need to be placed by experts, remain in the environment for 12 – 18 hours, are heavy, and require manual processing.



- Drone-based autonomous deployment for high-throughput

It takes too long to place traps. Ideal places to deploy traps are difficult for humans to reach. It make take several hours to place and collect a trap.

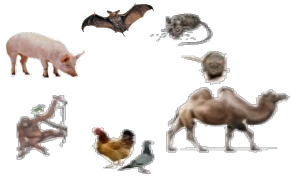


- Better metagenomics analytics to automatically identify threats

Today humans must sift through lots of biological data to pick out candidate threats. This would not scale for high volumes of field collections.

PREMONITION

High-throughput and low-cost monitoring known and unknown pathogens in the environment via autonomous collection of mosquitoes.



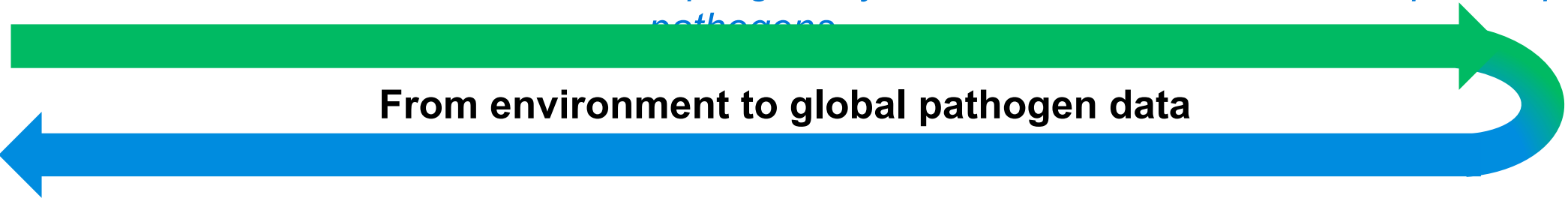
Animals maintain pathogens in environment



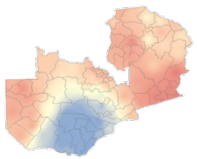
Mosquitoes collect blood, sampling many pathogens



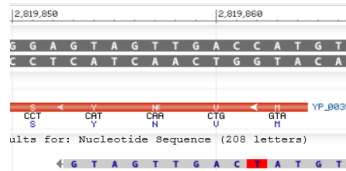
Drones deploy autonomous mosquito traps



From environment to global pathogen data



Global pathogen map of genes in space × time



Collected mosquitoes are gene sequenced and pathogens detected

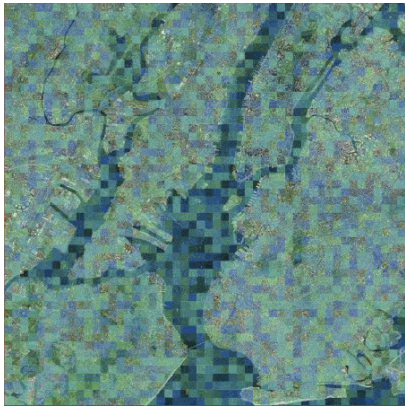


Long-range aerial or ground systems deploy short-range drones

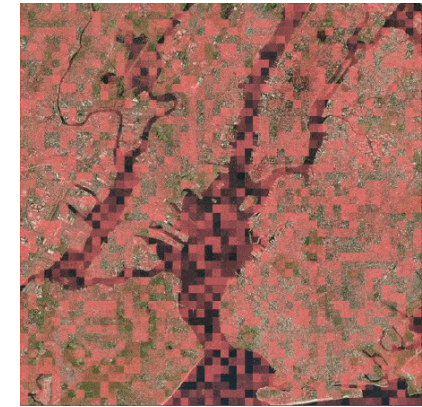
PREMONITION

Cloud-based genomic data analytics model epidemiology of emerging threats.

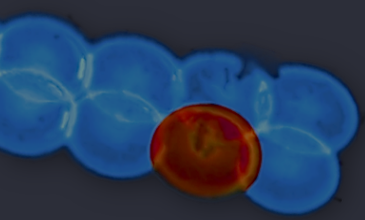
Example: Movement of potential pathogen and host observed through mosquitoes



Spatial/temporal
gene data of hosts
and vectors

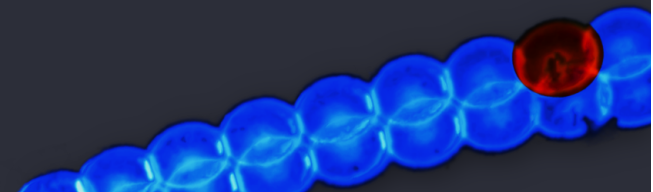


Spatial/temporal gene
data of potential
pathogens



The Grenada Experiment

Data and requirements from the
field



The Team



Amy Baldwin

St. George's University
Department of Microbiology



Jonathan Carlson

Microsoft Research
Computational Genomics



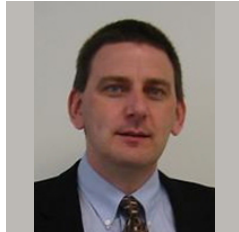
Ethan Jackson

Microsoft Research
Research in Software
Engineering Group



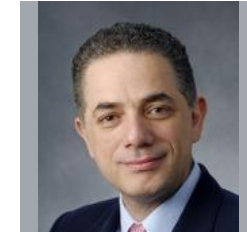
Ashish Kapoor

Microsoft Research
Adaptive Systems and
Interaction Group



Eamonn Keogh

University of California
Riverside
Department of Computer
Science and Engineering



Shawn Keshmiri

University of Kansas
Department of Aerospace
Engineering



Vijay Kumar

University of Pennsylvania
GRASP Laboratory



Douglas Norris

Johns Hopkins University
Department of Microbiology
and Immunology



James Pipas

University of Pittsburgh
Department of Molecular
Biology

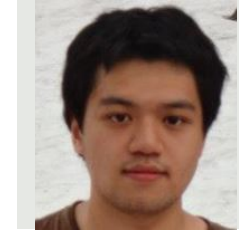
The Team



Shaz Qadeer
Microsoft Research
Research in Software
Engineering Group



Anandansankar Ray
University of California
Riverside
Department of Entomology



Alex Ching
Microsoft Research
Hardware Laboratory



Janos Sztipanovits
Vanderbilt University
Institute for Software
Integrated Systems



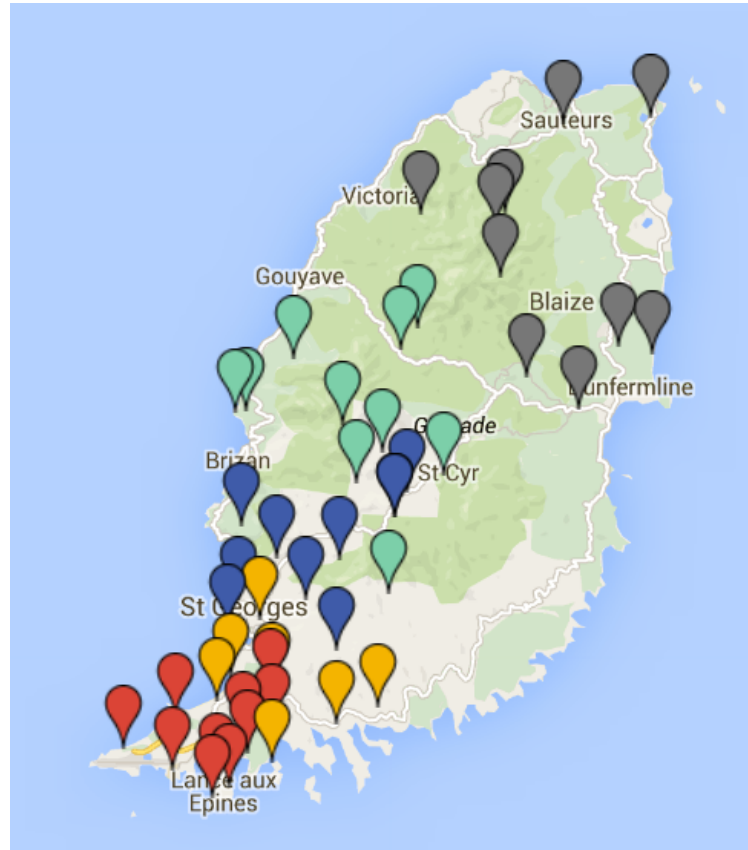
Michael Zyskowski
Microsoft Research
Outreach



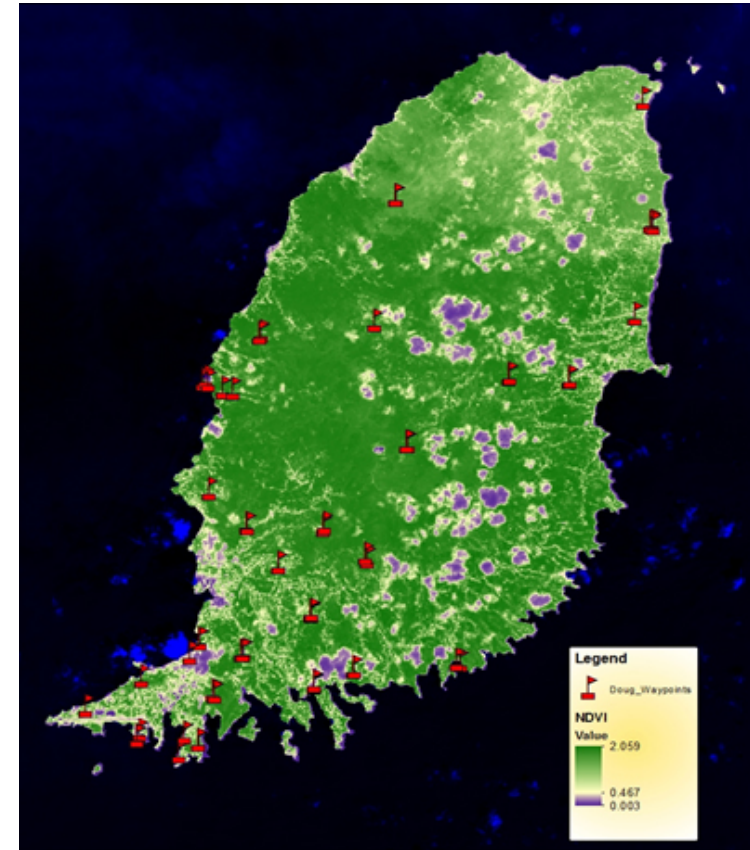
Patrick Therien
Microsoft Research
Hardware Laboratory

Sampling Grenada

Manual pilot in Grenada to understand feasibility and engineering constraints.
50 collections at 30 sites in urban, peri-urban, beach, and forest habitats



Planned



Actual (overlaid on 30m NDVI satellite data, courtesy Hopkins School of Public



Protocol – CO2 Baited CDC UV Trap

Placed during the day and collected the following morning.
Time in field: ~18 hours.



Urban



Beach

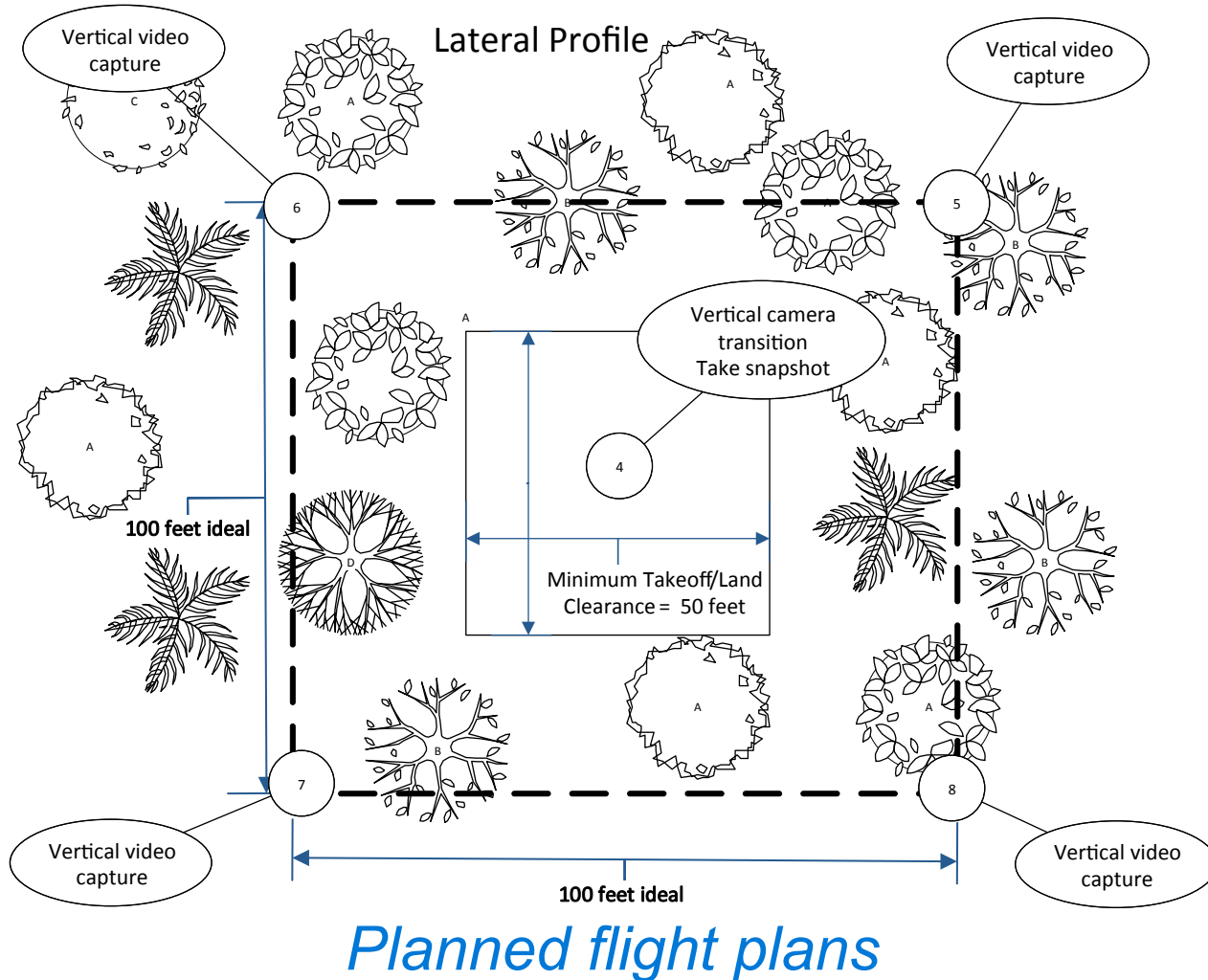


Jungle



Protocol – Drone Imagery

Manually collected drone imagery around the site to build mosquito hotspot classifiers, tree line detectors, and landing site detectors



Actual flights



Protocol – Sorting

Samples sorted by genus; subset of *Culex* and *Deinocerites* sorted by morphology.





St. George's University

DNA EXTRACTION

DNA EXTRACTION FOLLOWED BY PHOSPHORIC DNA EXTRACTION

1. Label three clean 1.5 ml microcentrifuge tubes.

2. Add 100 µl of lysis buffer to each tube.

3. Add 10 µl of DNA sample to each tube.

4. Vortex for 15 seconds.

5. Add 100 µl of 100% ethanol to each tube.

6. Vortex for 15 seconds.

7. Centrifuge at 14,000 rpm for 15 minutes.

8. Remove the supernatant.

9. Add 100 µl of 70% ethanol to each tube.

10. Vortex for 15 seconds.

11. Centrifuge at 14,000 rpm for 15 minutes.

12. Remove the supernatant.

13. Add 100 µl of 100% ethanol to each tube.

14. Vortex for 15 seconds.

15. Centrifuge at 14,000 rpm for 15 minutes.

16. Remove the supernatant.

17. Add 100 µl of 100% ethanol to each tube.

18. Vortex for 15 seconds.

19. Centrifuge at 14,000 rpm for 15 minutes.

20. Remove the supernatant.

21. Add 100 µl of 100% ethanol to each tube.

22. Vortex for 15 seconds.

23. Centrifuge at 14,000 rpm for 15 minutes.

24. Remove the supernatant.

25. Add 100 µl of 100% ethanol to each tube.

26. Vortex for 15 seconds.

27. Centrifuge at 14,000 rpm for 15 minutes.

28. Remove the supernatant.

29. Add 100 µl of 100% ethanol to each tube.

30. Vortex for 15 seconds.

31. Centrifuge at 14,000 rpm for 15 minutes.

32. Remove the supernatant.

33. Add 100 µl of 100% ethanol to each tube.

34. Vortex for 15 seconds.

35. Centrifuge at 14,000 rpm for 15 minutes.

36. Remove the supernatant.

37. Add 100 µl of 100% ethanol to each tube.

38. Vortex for 15 seconds.

39. Centrifuge at 14,000 rpm for 15 minutes.

40. Remove the supernatant.

41. Add 100 µl of 100% ethanol to each tube.

42. Vortex for 15 seconds.

43. Centrifuge at 14,000 rpm for 15 minutes.

44. Remove the supernatant.

45. Add 100 µl of 100% ethanol to each tube.

46. Vortex for 15 seconds.

47. Centrifuge at 14,000 rpm for 15 minutes.

48. Remove the supernatant.

49. Add 100 µl of 100% ethanol to each tube.

50. Vortex for 15 seconds.

51. Centrifuge at 14,000 rpm for 15 minutes.

52. Remove the supernatant.

53. Add 100 µl of 100% ethanol to each tube.

54. Vortex for 15 seconds.

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57. Add 100 µl of 100% ethanol to each tube.

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80. Remove the supernatant.

81. Add 100 µl of 100% ethanol to each tube.

82. Vortex for 15 seconds.

83. Centrifuge at 14,000 rpm for 15 minutes.

84. Remove the supernatant.

85. Add 100 µl of 100% ethanol to each tube.

86. Vortex for 15 seconds.

87. Centrifuge at 14,000 rpm for 15 minutes.

88. Remove the supernatant.

89. Add 100 µl of 100% ethanol to each tube.

90. Vortex for 15 seconds.

91. Centrifuge at 14,000 rpm for 15 minutes.

92. Remove the supernatant.

93. Add 100 µl of 100% ethanol to each tube.

94. Vortex for 15 seconds.

95. Centrifuge at 14,000 rpm for 15 minutes.

96. Remove the supernatant.

97. Add 100 µl of 100% ethanol to each tube.

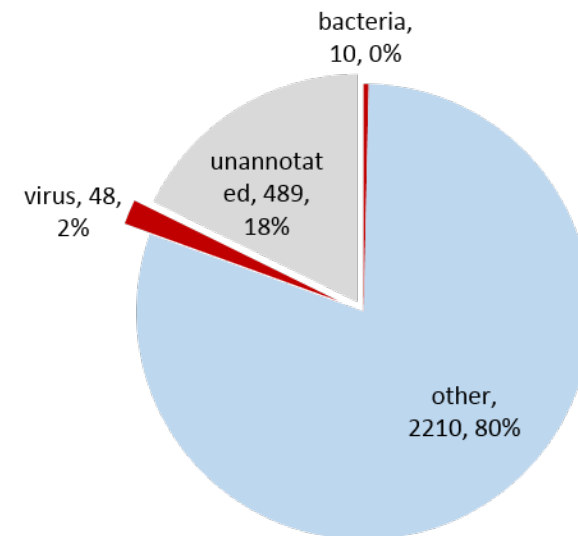
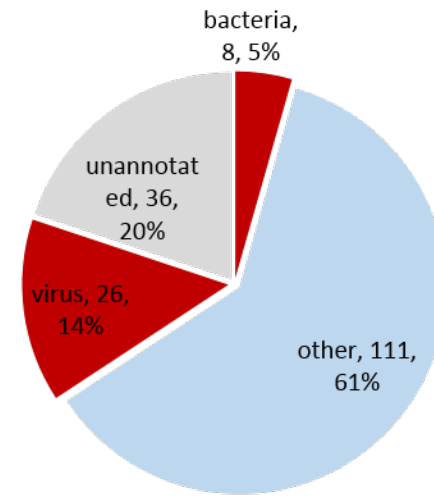
98. Vortex for 15 seconds.

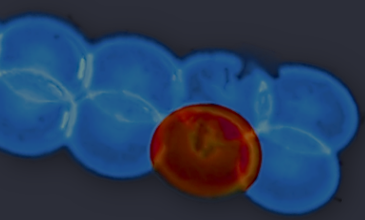
99. Centrifuge at 14,000 rpm for 15 minutes.

100. Remove the supernatant.

Protocol – Nucleic Acid Extraction

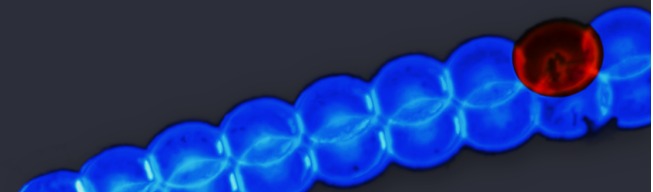
DNA/RNA aliquots deep sequenced to be computationally analyzed.





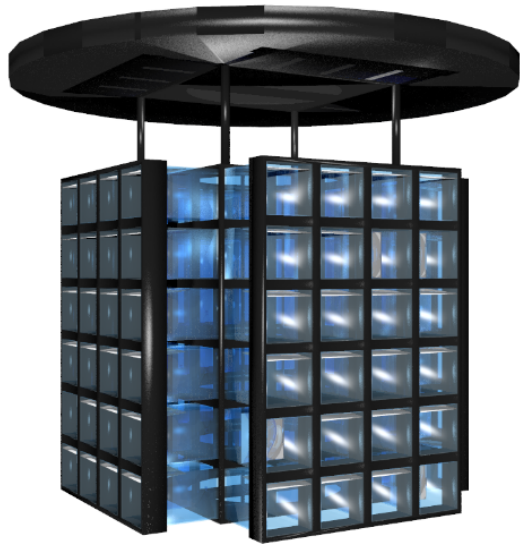
Building PREMONITION

Autonomous Traps

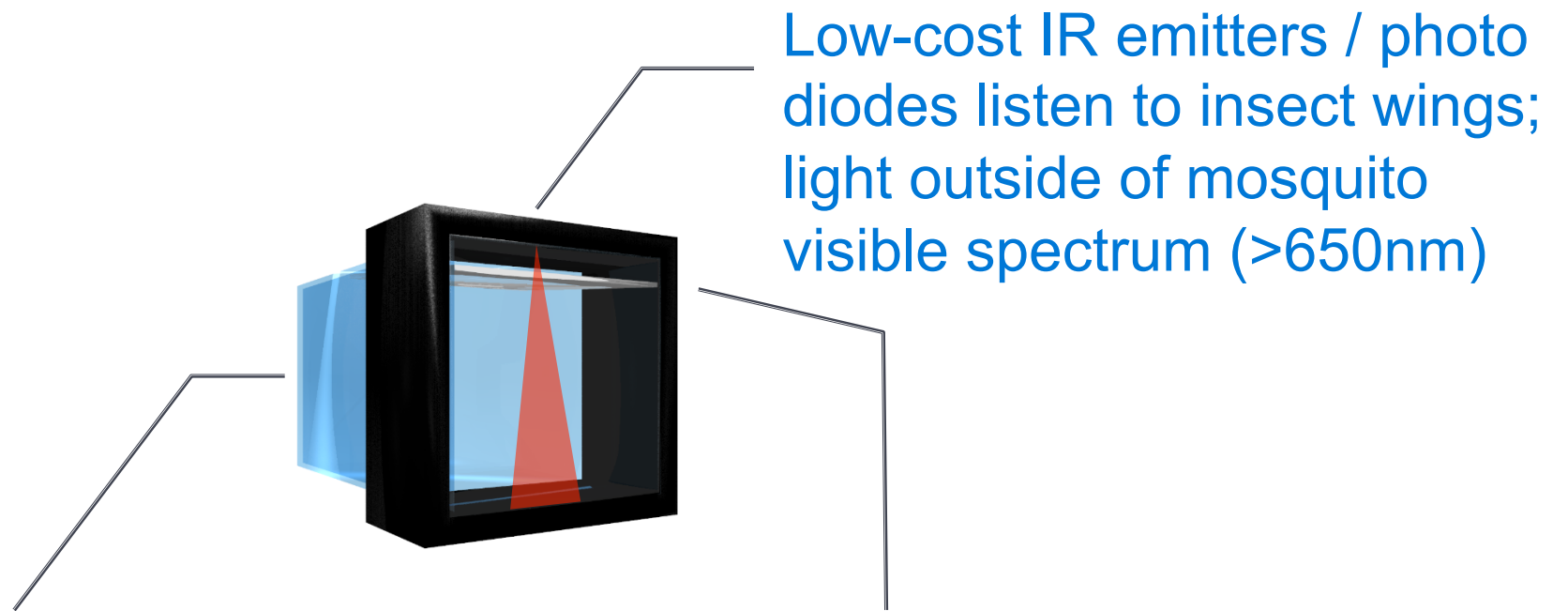


A Smarter Trap

Real-time speciation and selective trapping of insects; in-situ stabilization of nucleic acids; low-cost and low-power sensor and actuators.



Honeycomb design simplifies automated processing

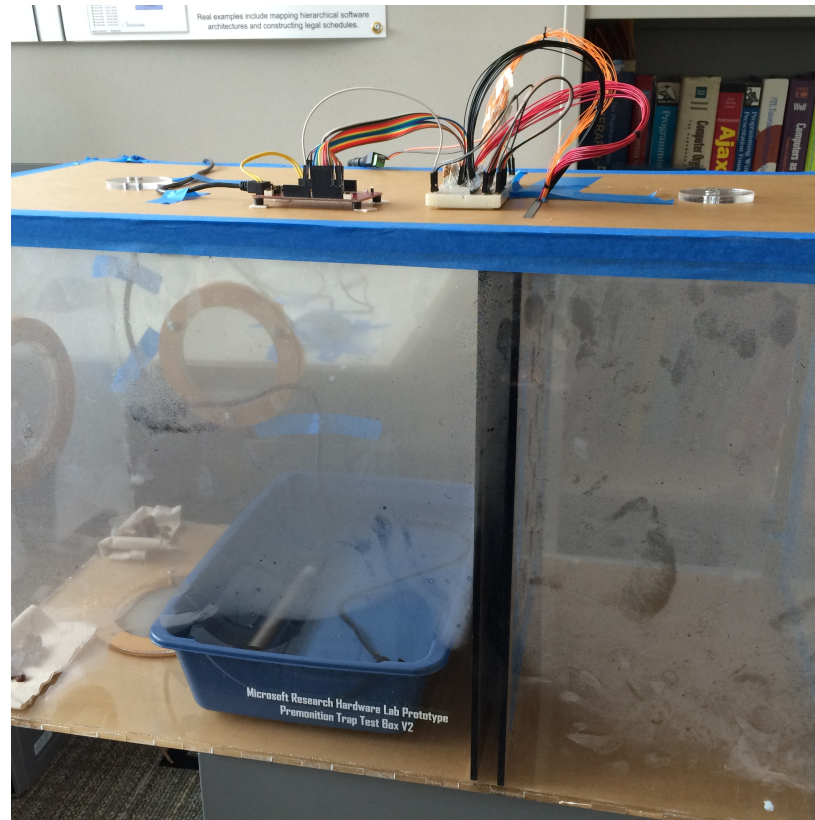
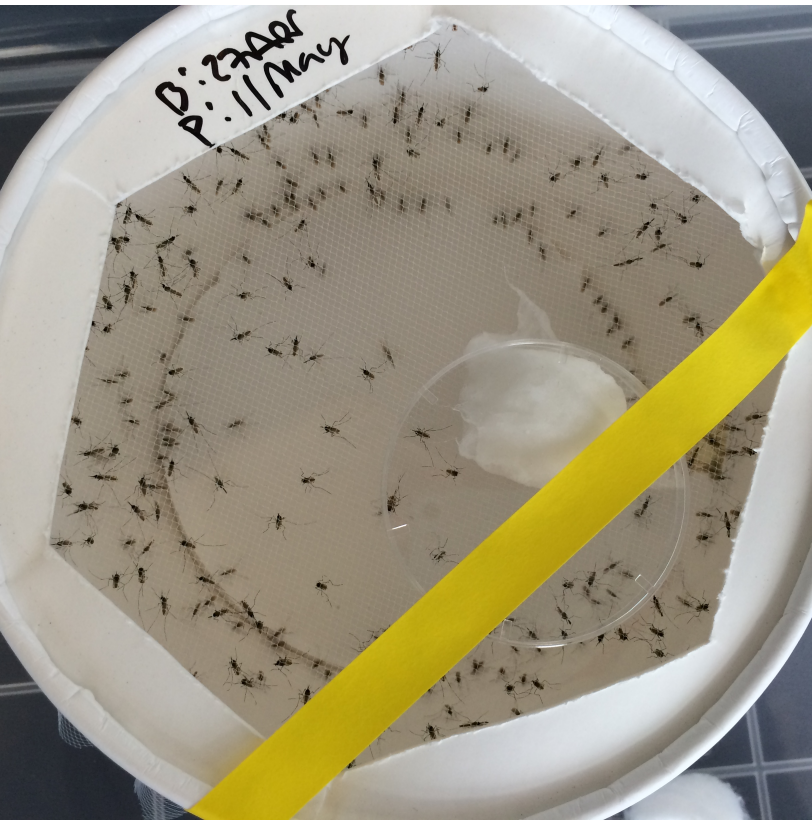


In-situ chemical stabilization of insect preserves sample for up to one week

Door closes when species of interest enters cell. Signal processing delay ~100ms

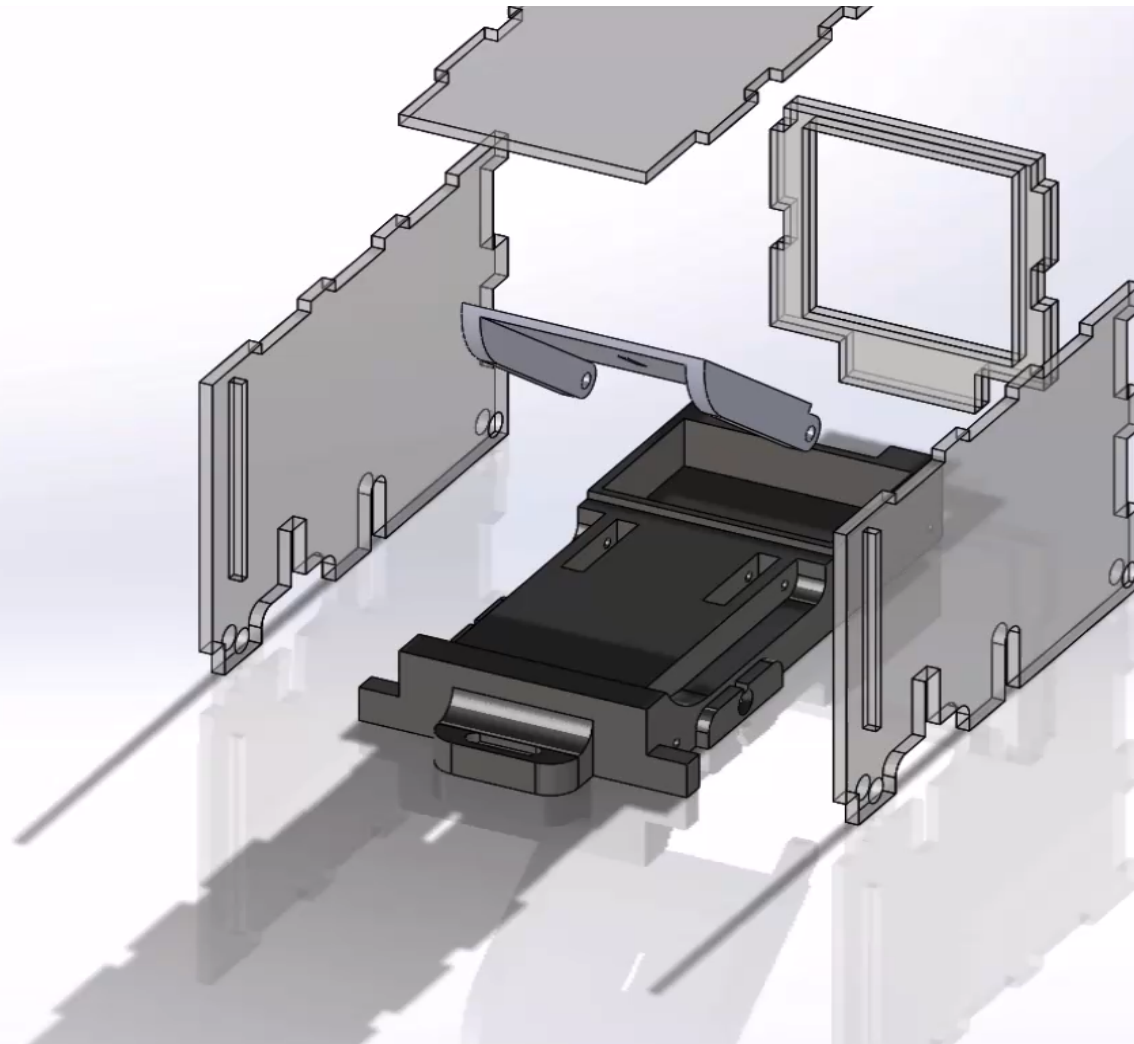
Designed and tested sensors and actuators

In laboratory setting on lab-raised anopheles (malaria) mosquitoes



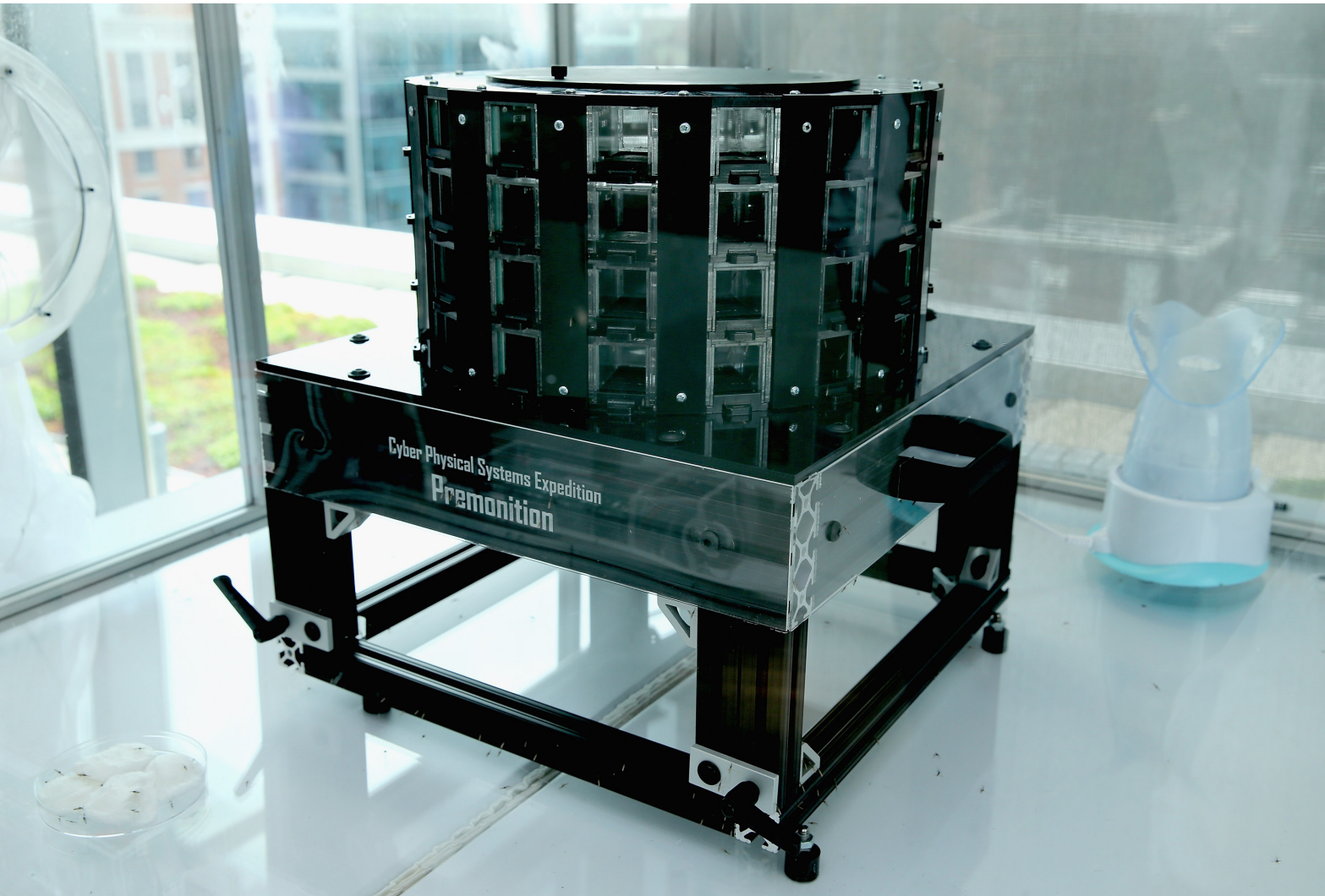
A Challenging CPS Design Problem

Co-design problem involving: cost, manufacturability, harsh environments, computational power, energy constraints, and size and weight

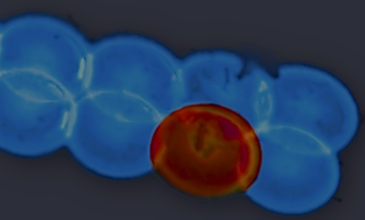


Most Advanced Mosquito Trap in World

Real-time detection and preservation using low-cost sensors, embedded software, and machine learning

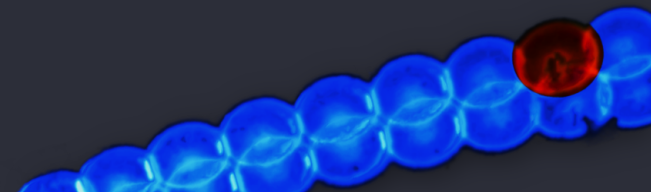


Cells have no electronics or motors



Building PREMONITION

Autonomous Deployment



Autonomous Deployment

Developing a drone control stack using advanced operating systems, program verification technologies, vision, and machine learning

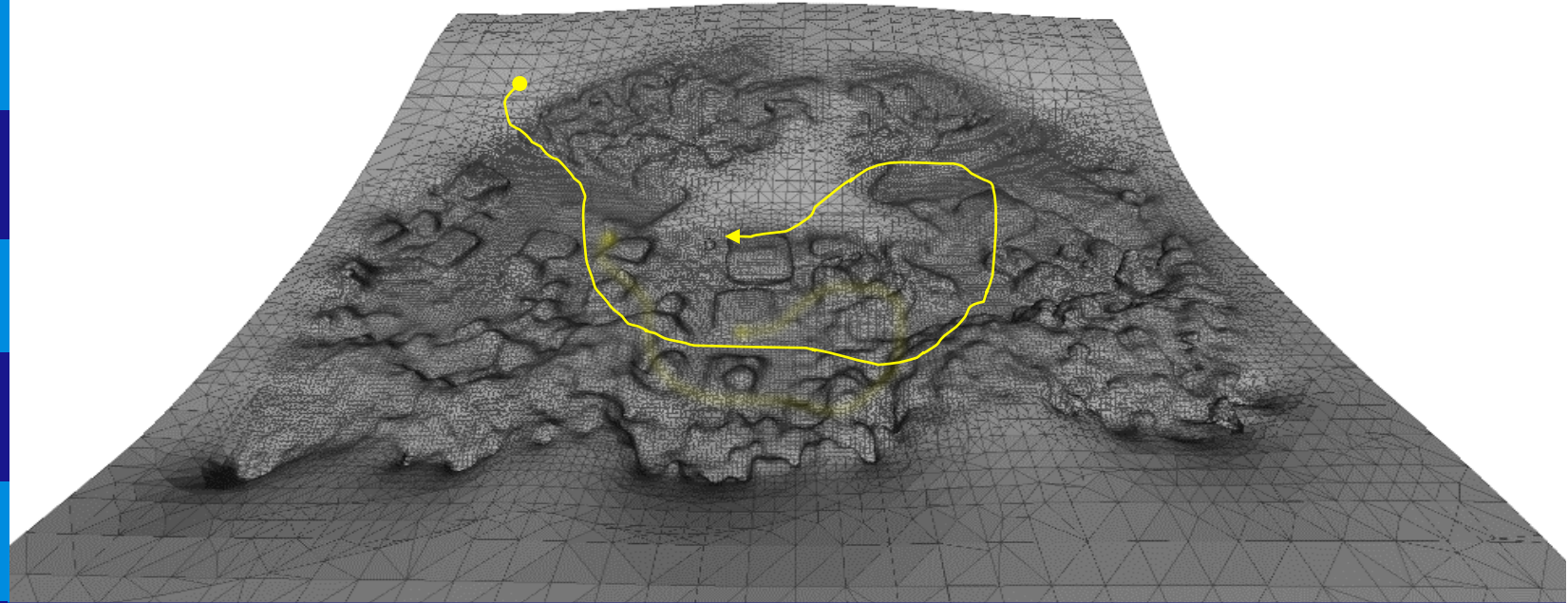
High-level Planning

Safe despite limited power, external disturbances,

Correct Control

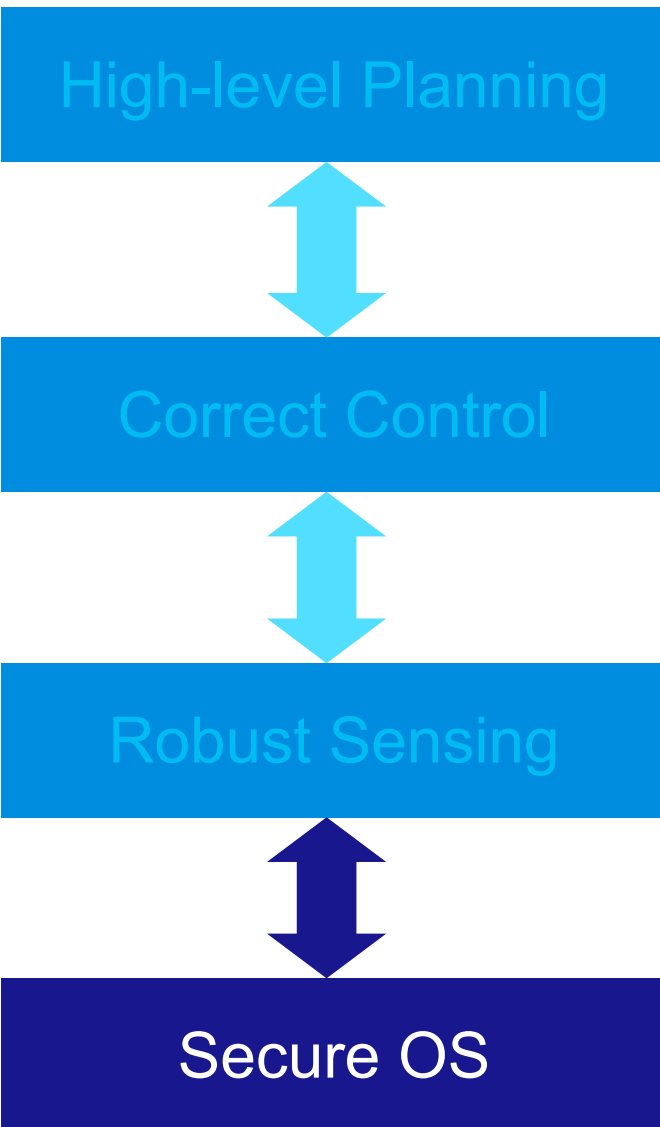
Robust Sensing

Secure OS



sensor noise, and complex missions

Autonomous Deployment



Drone OS:

Provides secure and robust infrastructure for embedded software

Guarantees:

Full functional correctness of OS components and protocols

Methods:

Built from Verve OS; mechanically verified using Z3, Boogie

Verified
**Garbage
Collector**

Verified
Threads

Verified
**Interrupt
Handlers**

Verified
**Device
Interface**

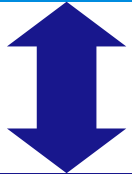
Verified
Startup

Autonomous Deployment

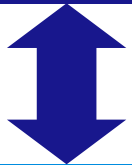
High-level Planning



Correct Control



Robust Sensing



Secure OS

Drone Vision:

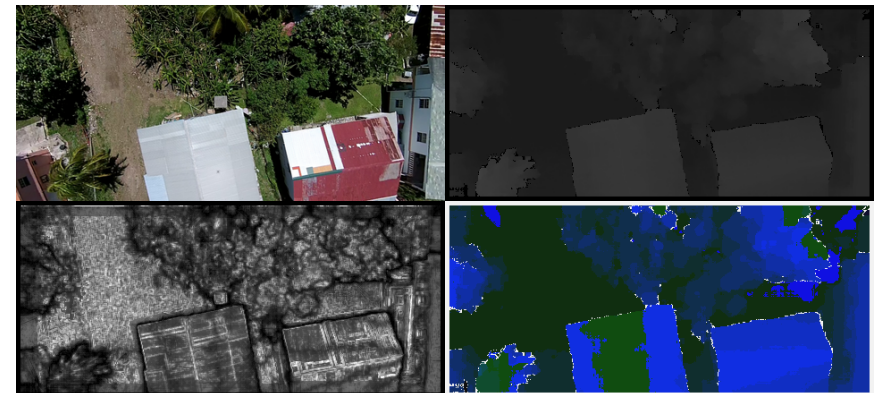
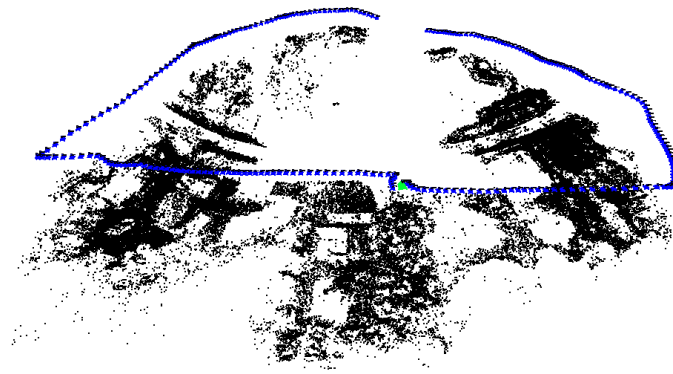
Reconstructs 3D environment and camera poses

Guarantees:

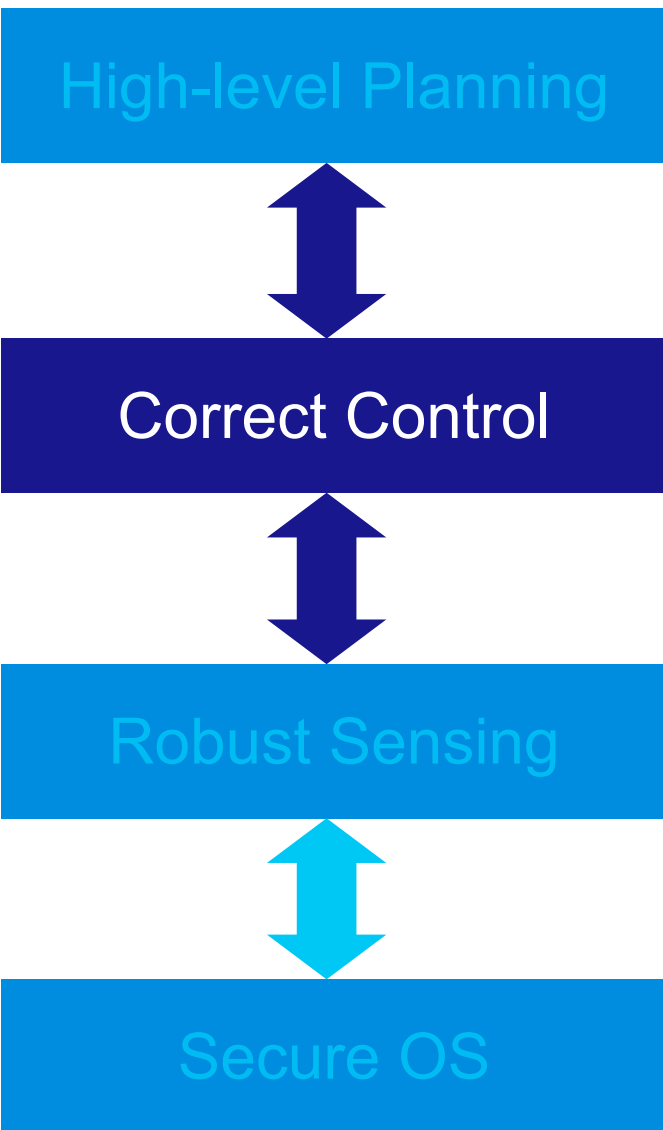
Goodness of classifications (Bayesian posterior intervals)

Methods:

Built from MSR vision libs and Bayesian ML



Autonomous Deployment



Controller synthesis and validation:

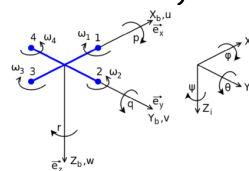
Designs safe control laws handling probabilistic sensor data

Guarantees:

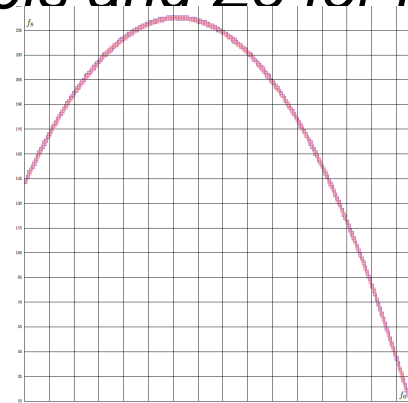
Satisfaction of (probabilistic) safety constraints – extensions of LTL

Methods:

Optimal control, Taylor models and Z3 for hybrid validation

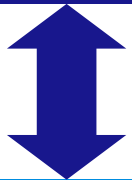


$$\begin{aligned}
 \dot{p} &= \frac{l_x}{l_y} \sum_{i=0}^3 \gamma_i (\omega_2^i - \omega_4^i) + \frac{l_x}{l_y} q \sum_{j=1}^4 \omega_j (-1)^j + \frac{l_y - l_x}{l_y} qr \\
 \dot{q} &= \frac{l_x}{l_y} \sum_{i=0}^3 \gamma_i (\omega_3^i - \omega_1^i) + \frac{l_x}{l_y} q \sum_{j=1}^4 \omega_j (-1)^j + \frac{l_x - l_y}{l_y} pr \\
 \dot{r} &= \frac{1}{l_y} \sum_{j=1}^4 (l_c \dot{\omega}_j + k_D \omega_j^2 + B_a \omega_j) (-1)^j \\
 \dot{u} &= vr - wq - g \sin(\theta) \\
 \dot{v} &= wp - ur + g \sin(\phi) \cos(\theta) \\
 \dot{w} &= uq - vp + g \cos(\phi) \cos(\theta) - \frac{1}{m} \sum_{j=1}^4 \sum_{i=0}^3 \gamma_i \omega_j^i
 \end{aligned}$$



Autonomous Deployment

High-level Planning



Correct Control



Robust Sensing



Secure OS

High-level PL for planners:

DSL for asynchronous event-driven programming of planners

Guarantees:

Safety and liveness properties over event streams

Methods:

P language, Zing model checker, FORMULA DSL tools

Motion Planner

Plan Executor

Sense and Infer

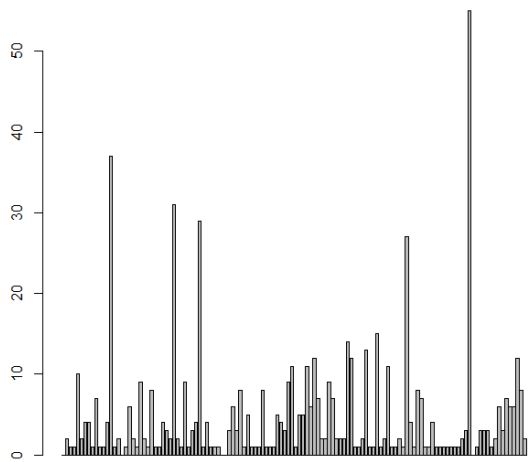
State Manager

P code

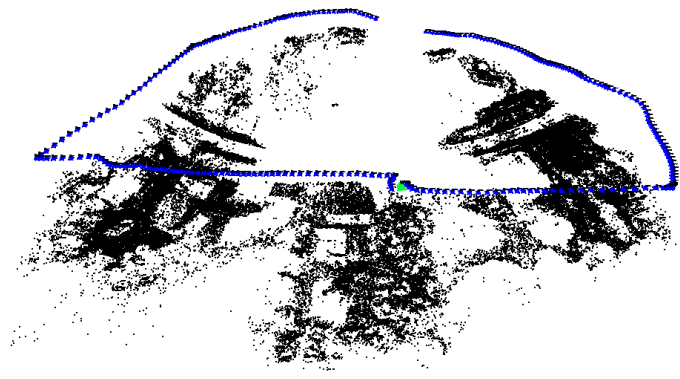


Learning Mosquito Hotspots

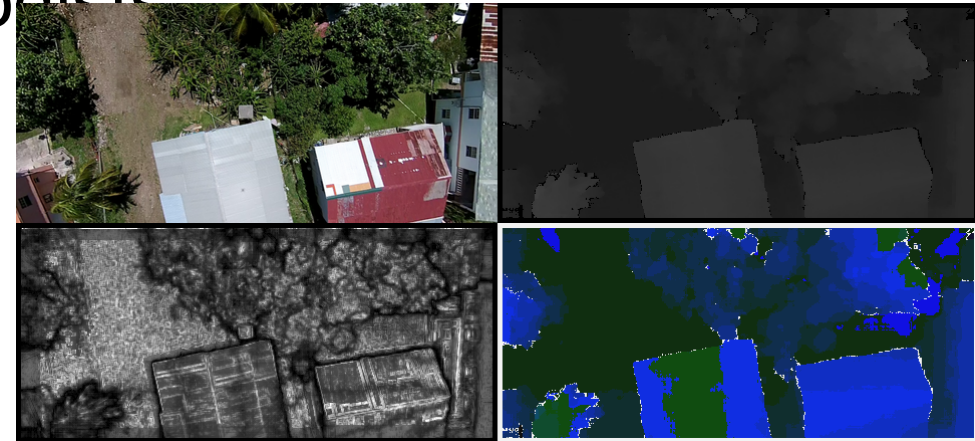
So drones can make deployment decisions that normally require field biologists



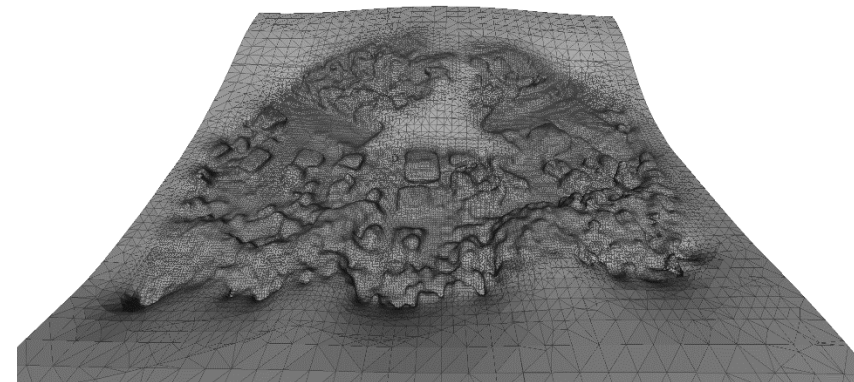
Mosquito counts by site



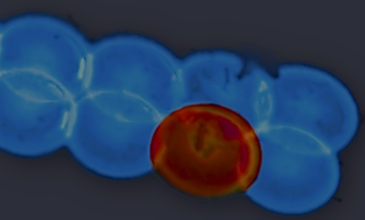
*Reconstruction of camera pose
and drone path*



Mosquito hotspot

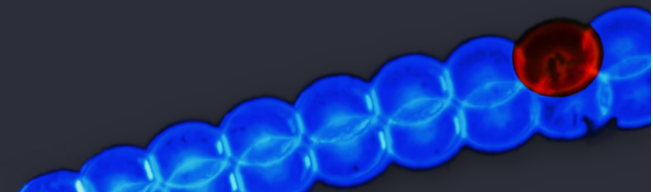


*Reconstruction of fined-grained
3D environment*



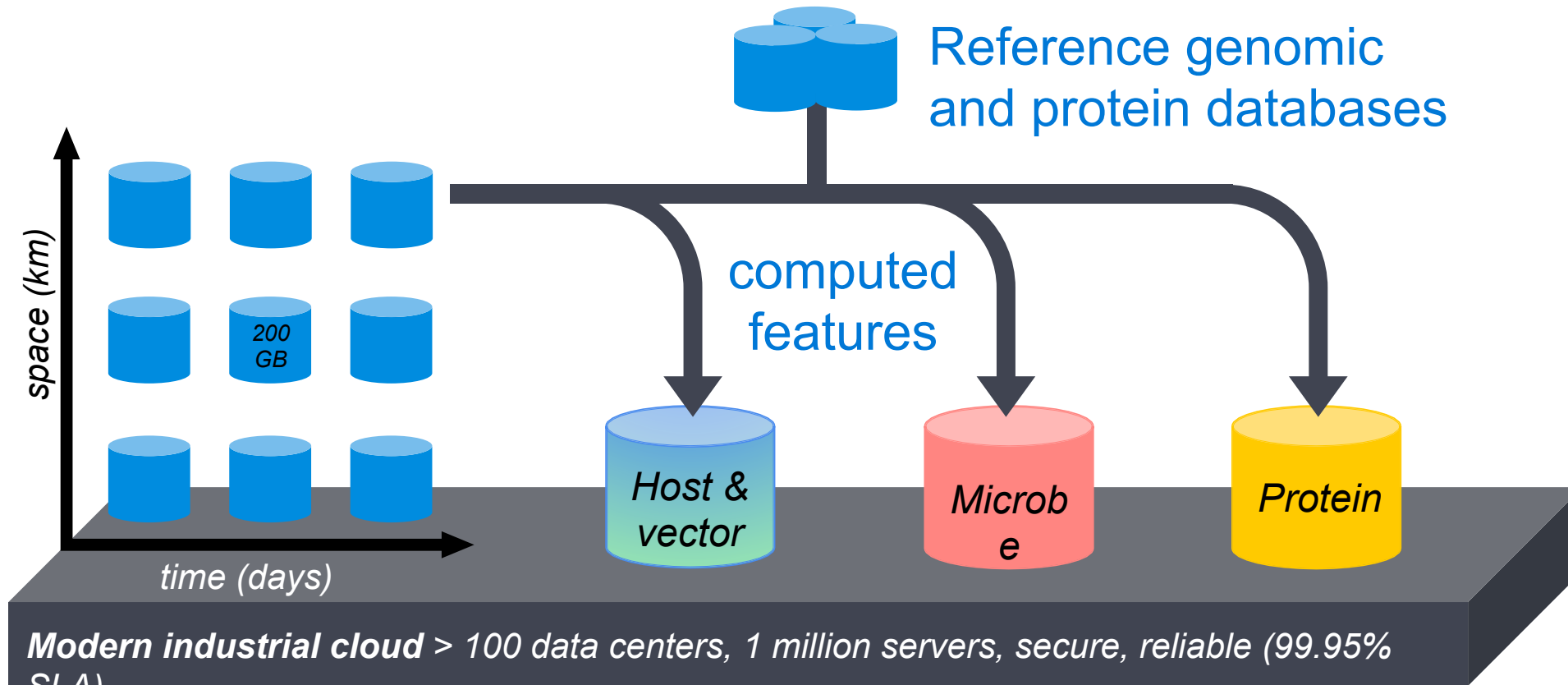
Building PREMONITION

Autonomous threat detection



Better genomic data analytics

Feature extraction and indexing from spatial/temporal genetic data.



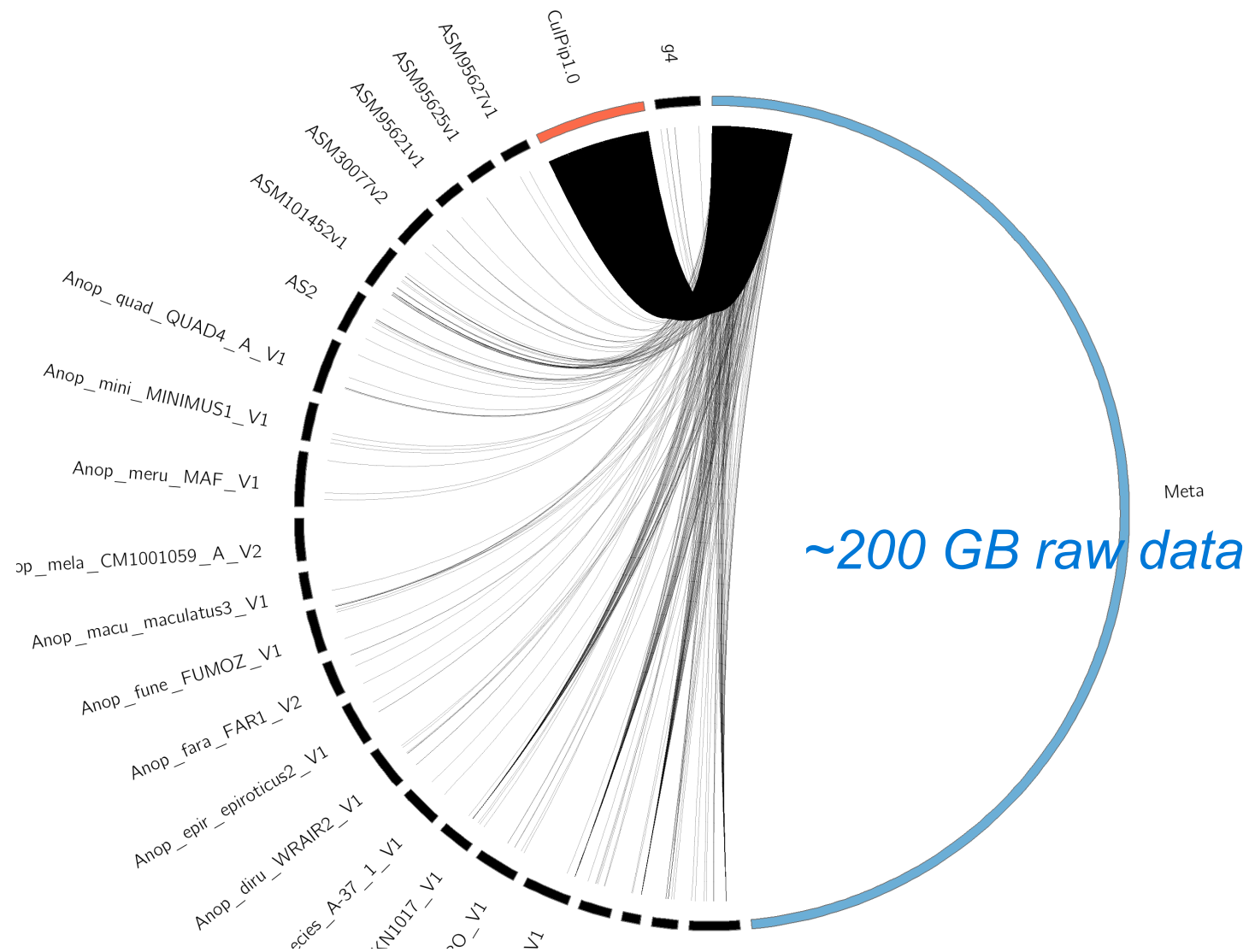
Data cleansing and similarity to reference genes

De novo assembly to reconstruct genes of unknown organisms

Recognize potential threat sequences by expressed proteins

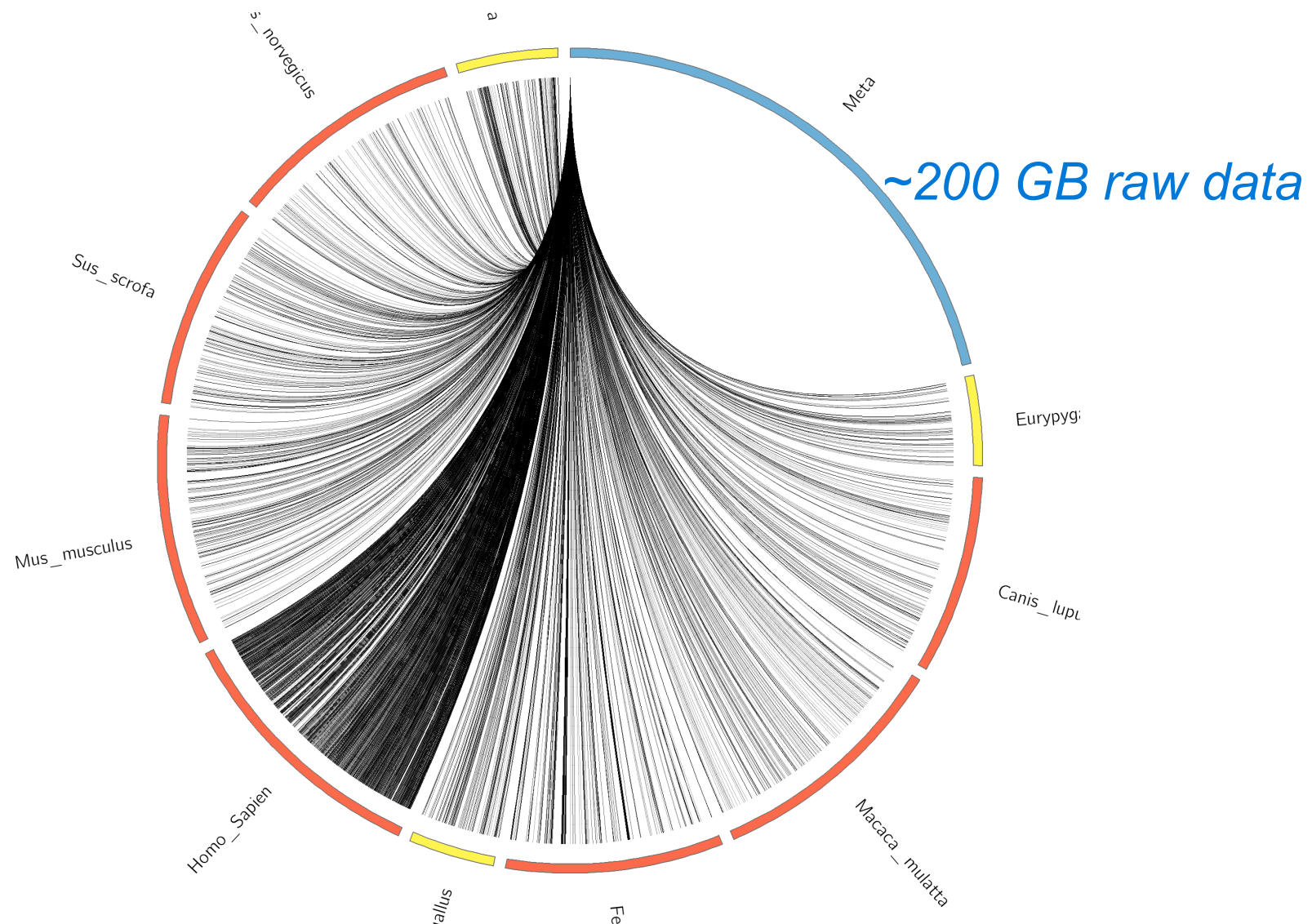
Vector data

We can recover the species of mosquitoes collected in Grenada



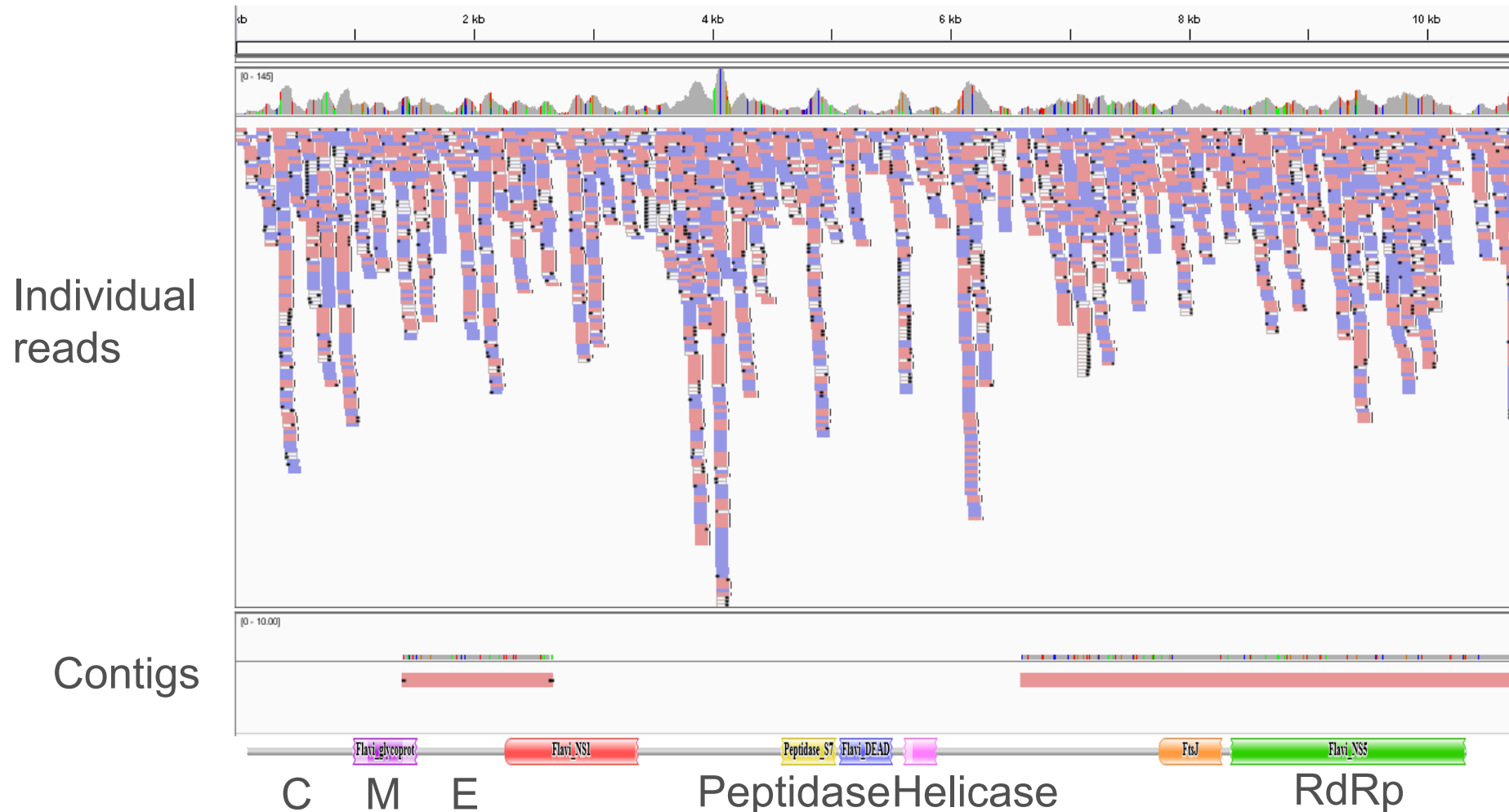
Host data

We can recover what mosquitoes bit in Grenada

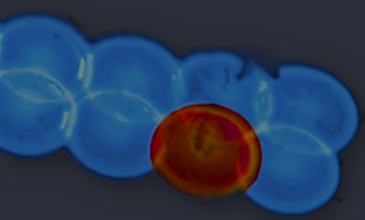


Virus data

We can recover novel potential pathogens from mosquitoes in Grenada

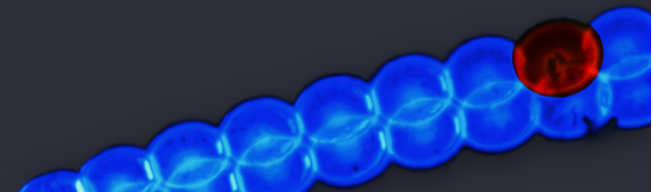


A computationally reconstructed candidate novel Culex Flavivirus



The Path Ahead

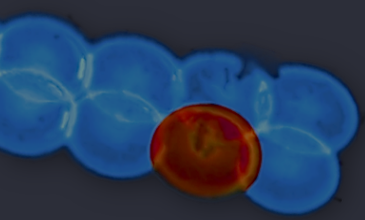
How far off is it?



How far off is it?

We estimate 5 years because the technology trends are clear:

- Moore's law on microprocessors means field entomology can be automated at scale.
- Commodity drones and robotic platforms are here to stay. Innovations in 3D printing continue to fuel a Cambrian explosion.
- Gene sequence costs decreasing faster than Moore's law, and cloud capacity continues to grow. Gene sequencing will enable high-throughput surveillance of (unknown) pathogens.



Thank You

Questions are welcome

