

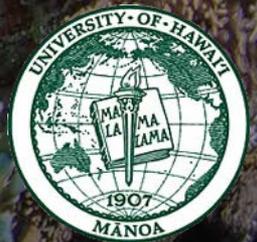
Where are all the fish going? Identifying patterns of genetic connectivity across the Hawaiian Archipelago

Richard R. Coleman, Ph.D

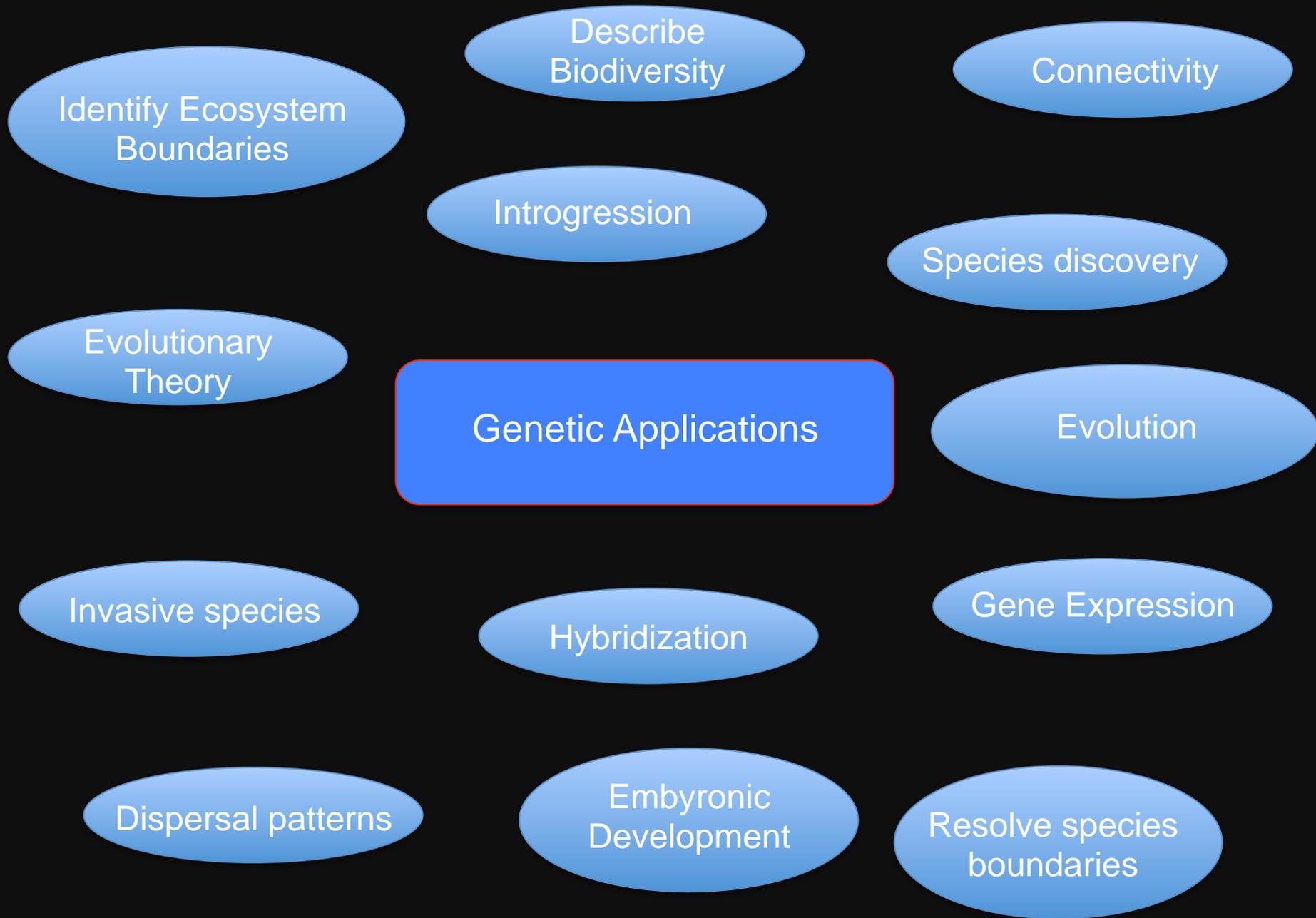
Hawai'i Institute of Marine Biology
University of Hawai'i, Mānoa, Dept of Biology

24 September 2019

National Marine Sanctuaries Webinar Series







Identify Ecosystem Boundaries

Describe Biodiversity

Connectivity

Introgression

Species discovery

Evolutionary Theory

Genetic Applications

Evolution

Invasive species

Hybridization

Gene Expression

Dispersal patterns

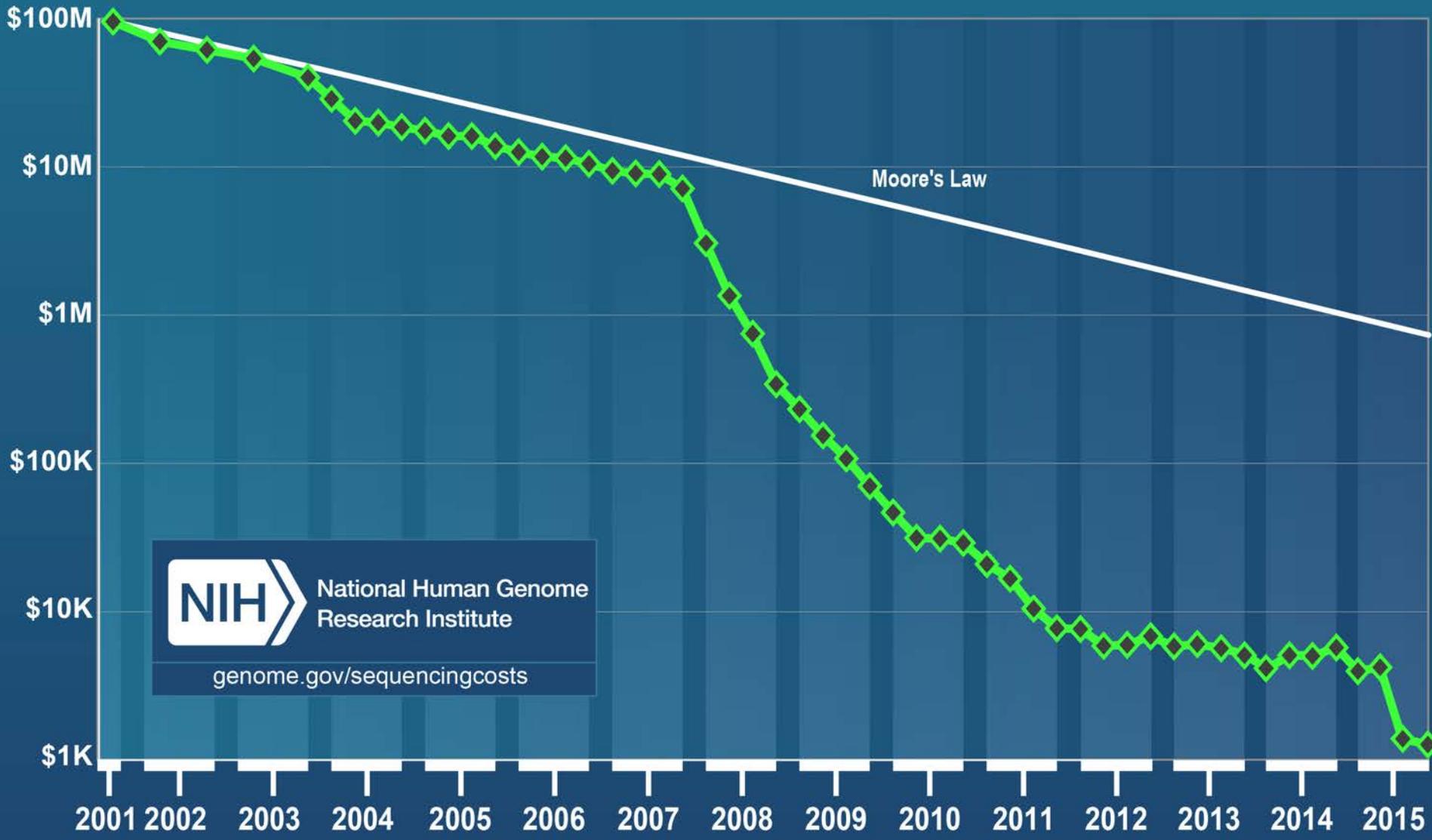
Embryonic Development

Resolve species boundaries

Brief History of Genetics

- 1859 – “On the origin of species” Theory of Evolution by Natural Selection
- 1865 – Gregor Mendel’s pea experiments
- 1869 – DNA isolated
- 1953 – DNA double helix described (Watson, Crick, Franklin, Wilkins)
- 1966 – Genetic code cracked
- 1975 – DNA Sequencing
- 1983 – PCR invented
- 1995 – First genome sequenced
- 2002 – First mammal genome sequenced
- 2003 – Human genome sequenced
- 2005 – Second generation sequencing (Roche 454)
- 2006 – Illumina sequencing (accounts for >70% of the market)

Cost per Genome



NIH National Human Genome Research Institute
genome.gov/sequencingcosts



Skillings et al. 2014



Timmers et al. 2012



Andrews et al. 2010



Ahti et al. 2016



Johnston et al. 2017



Waldrop et al. 2016



Gaither et al. 2013



Iacchei et al. 2015



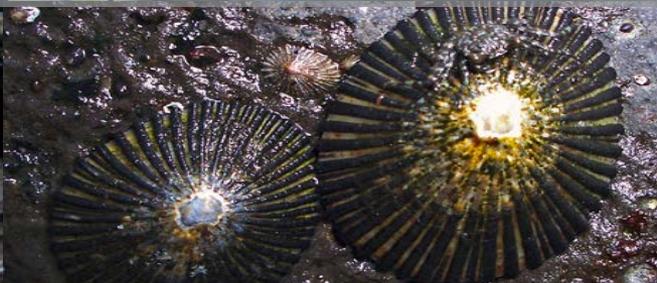
Fernandez-Silva et al. 2016



Puritz & Toonen 2011



DiBattista et al. 2013



Bird et al. 2013

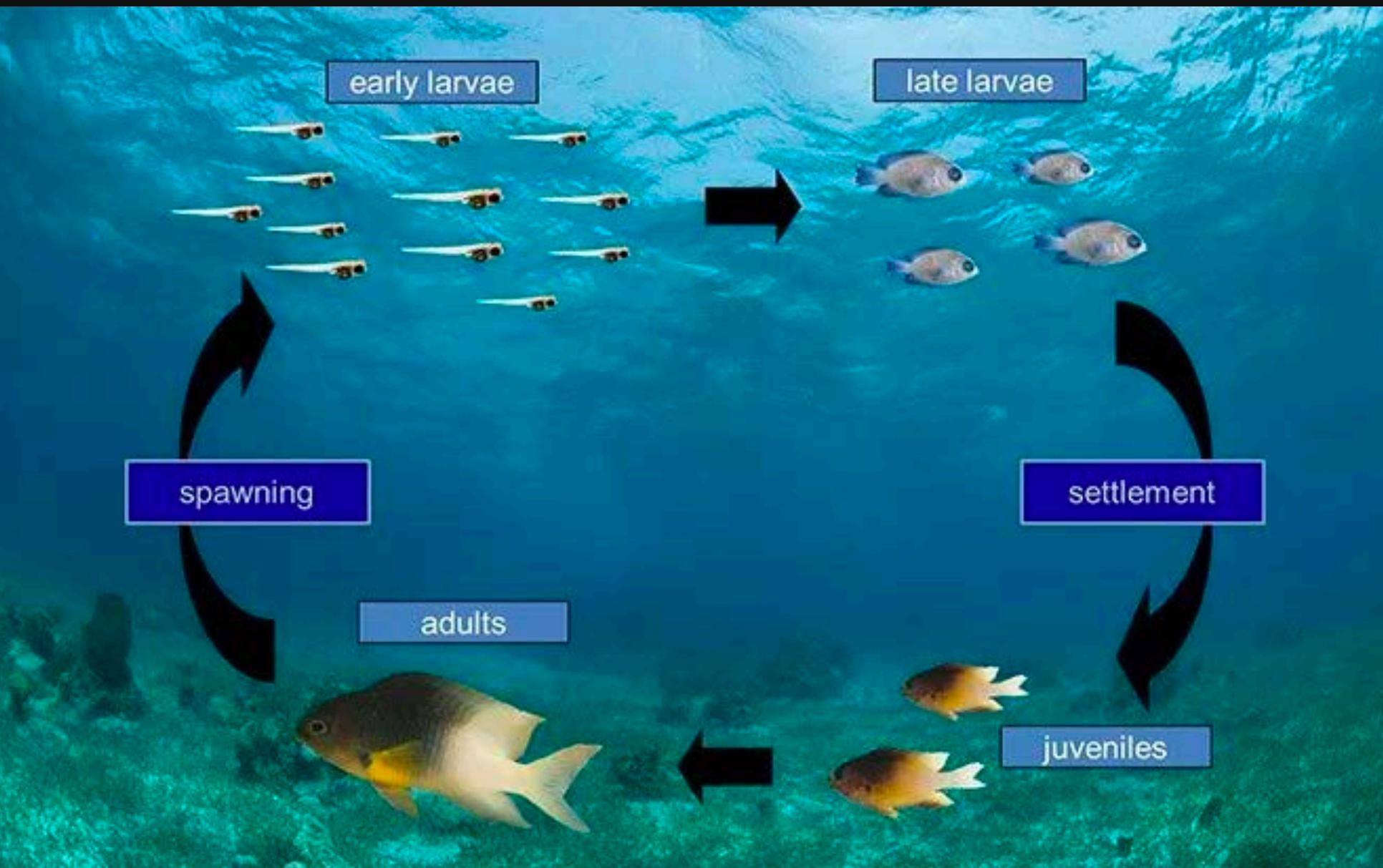


Importance of Understanding Connectivity

- Provides insights into the mechanisms that influence evolution
- Information can be used to inform management and conservation strategies

Informing proper management

- Barriers to dispersal
- Identify vulnerable areas and management units
- Characterize source-sink populations
 - “Source” populations
 - Populations that are responsible for seeding other areas
 - “Sink” populations
 - Regions that are dependent on other populations



early larvae

late larvae

40-60 day

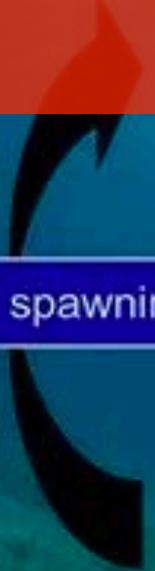


spawning

settlement

adults

juveniles



Population Connectivity



Population Connectivity



Population Connectivity



Population Connectivity



Population Connectivity



Population Connectivity



Population Connectivity



Barriers to dispersal in terrestrial systems



wallpaperswiki.com

- Often easy to identify
- Examples:
 - Rivers
 - Mountain Ranges
 - Islands

Barriers in marine systems



Most marine organisms have larvae that can drift for weeks to months



Spatial Scales



Spatial Scales



Ocean Basins

Range-wide (Indian and Pacific Oceans)

Spatial Scales



Ocean Basins

Range-wide (Indian and Pacific Oceans)

Archipelago

Hawaiian Archipelago

Spatial Scales



Ocean Basins

Range-wide (Indian and Pacific Oceans)

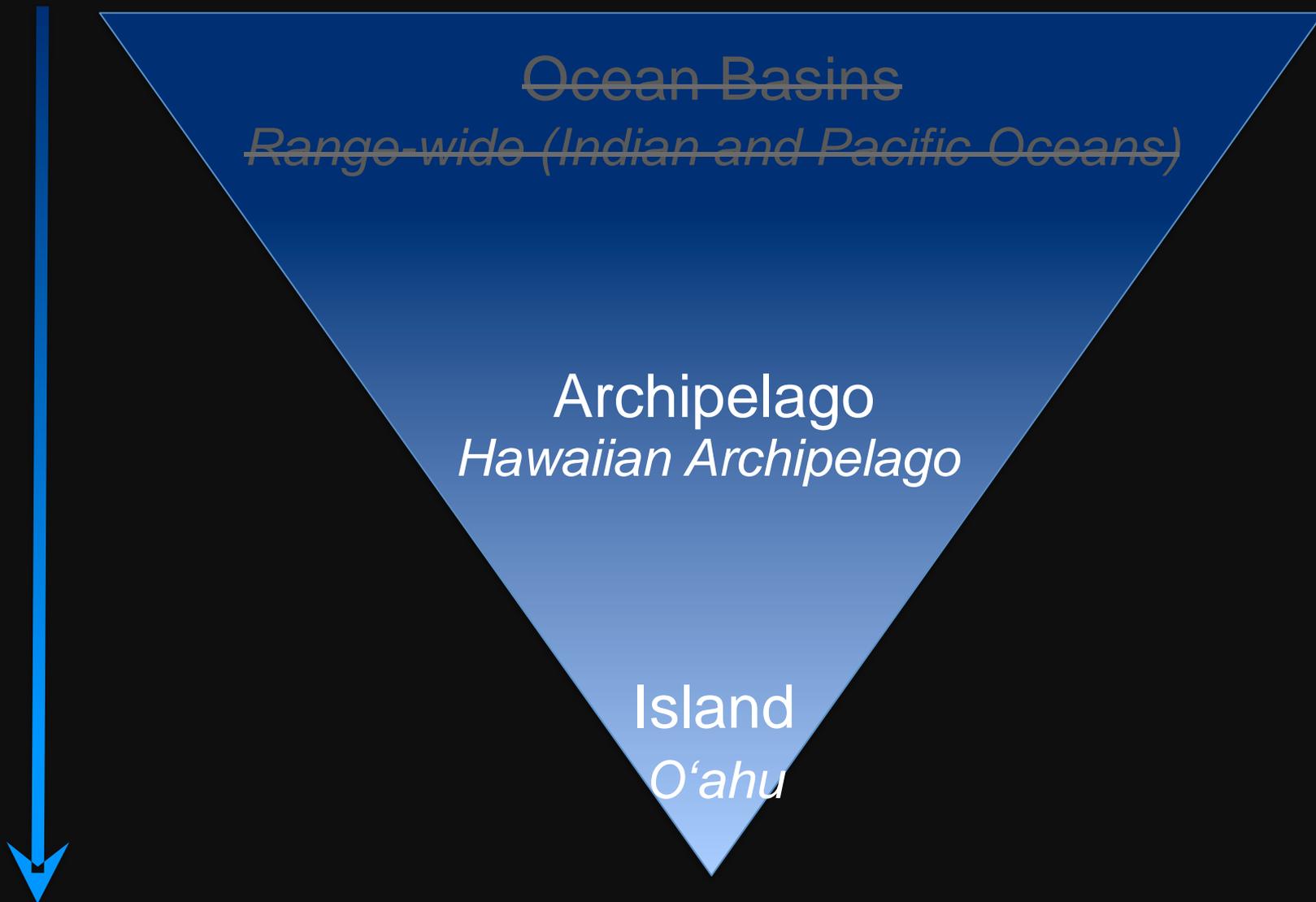
Archipelago

Hawaiian Archipelago

Island

O'ahu

Spatial Scales



Spatial Scales



Ocean Basins

Range-wide (Indian and Pacific Oceans)

Archipelago

Hawaiian Archipelago

Conservation/Management

Island

O'ahu



KAUAI
NIIHAU
Waialeale
5148 ft (1,569 m)

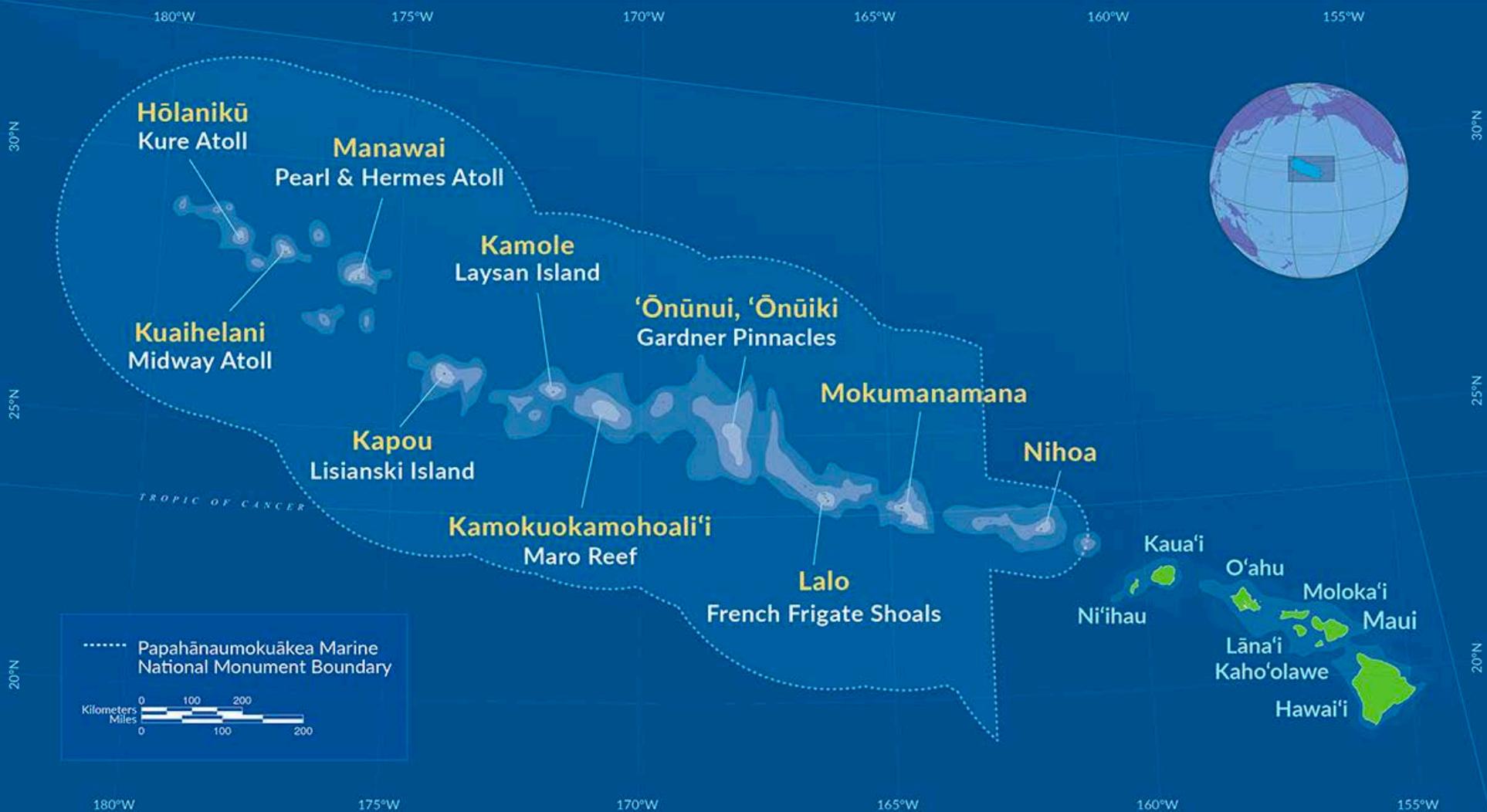
OAHU
Honolulu

MOLOKAI
LANAI

KAHOOLAWE
MAUI
Red Hill
10023 ft (3,055 m)

Upolu Pt
HAWAII
Mauna Kea
13796 ft (4,207 m)
Hilo
Cape Kumukahi
Mauna Loa
13677 ft (4,169 m)
Ka Lae

Hawaiian Archipelago



Main Hawaiian Islands



Midway Atoll



'Ōnūnui,
Gardner P

Mokumanamana



e Shoals



..... Papahānaumokuākea National Monument Boundary



30°N

25°N

20°N

30°N

25°N

20°N

170°W

165°W

160°W

155°W

180°W

175°W

170°W

165°W

160°W

155°W

Papahānaumokuākea Marine National Monument



Atoll
Kamole
Laysan Island



Mokumanamana



Lalo
French Frigate Shoals



ai



Archipelago

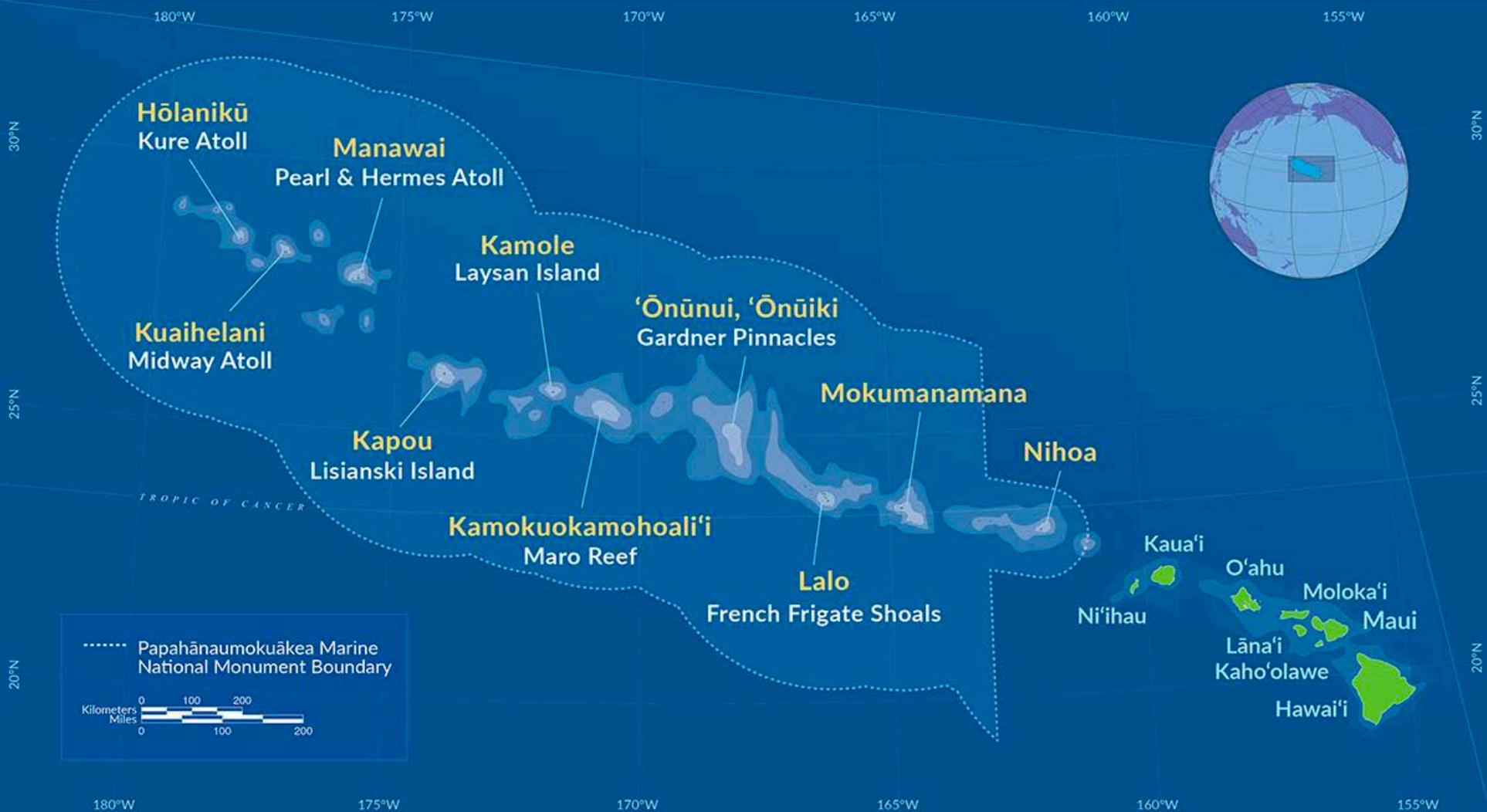
Island



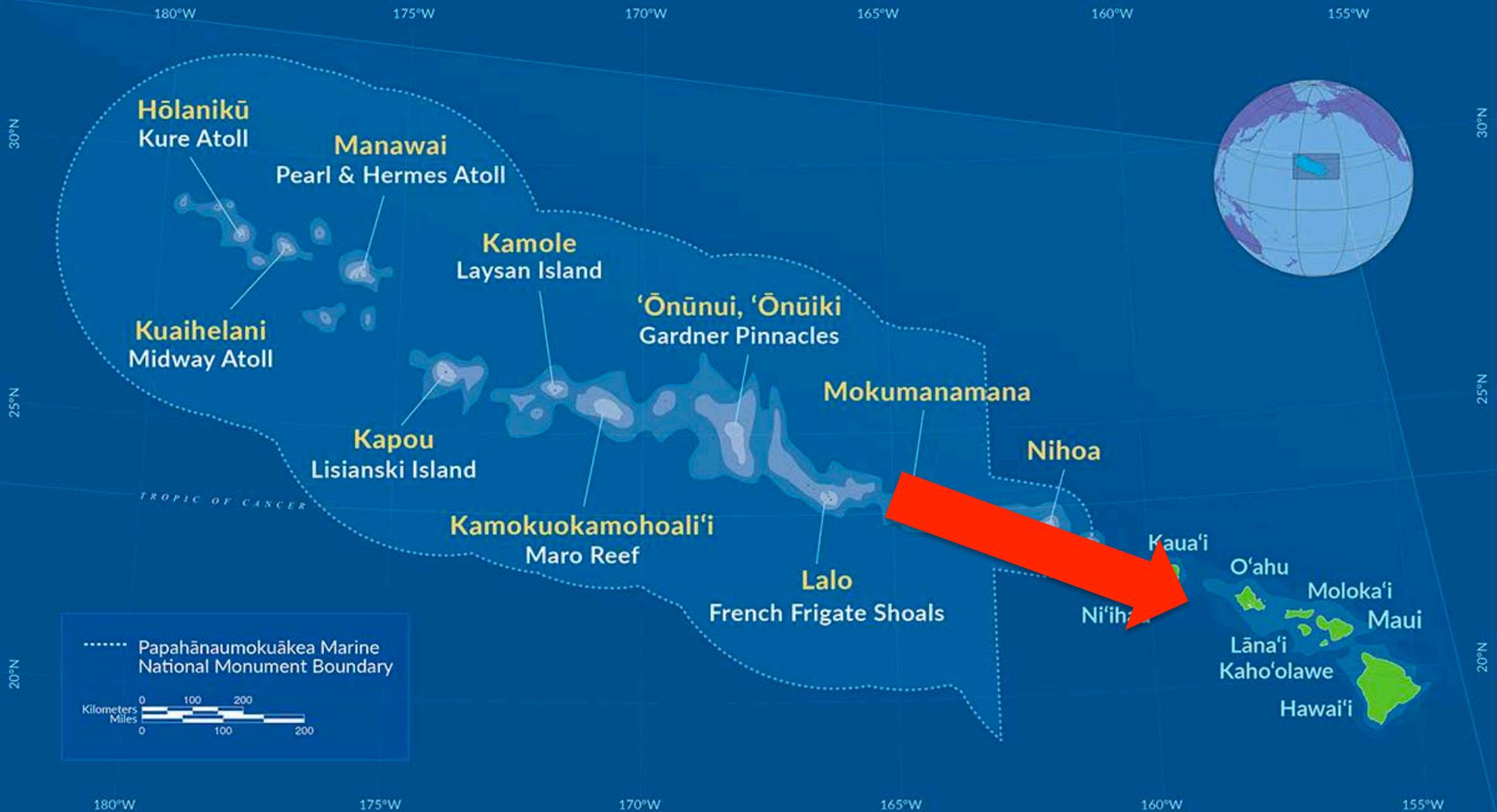


Map of the USA with the Hawaiian Archipelago overlaid. Black outline shows the boundary of the Northwestern Hawaiian Islands Coral Reef Ecosystem Reserve, which is 1200 Nautical Miles long by 100 Nautical Miles wide and has an area roughly the same size as Florida and Georgia combined, or 131,800 square statute miles.

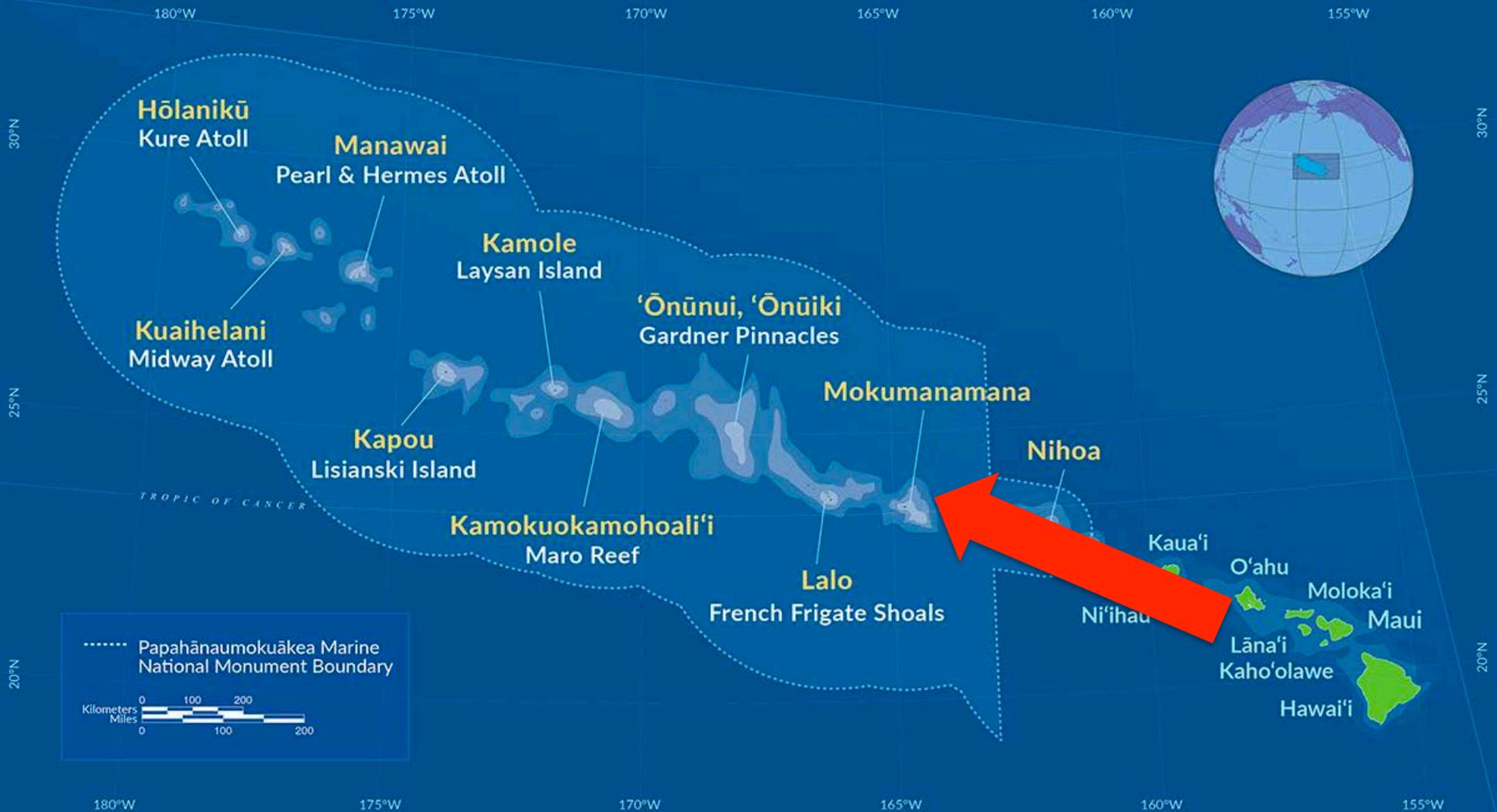
Hawaiian Archipelago



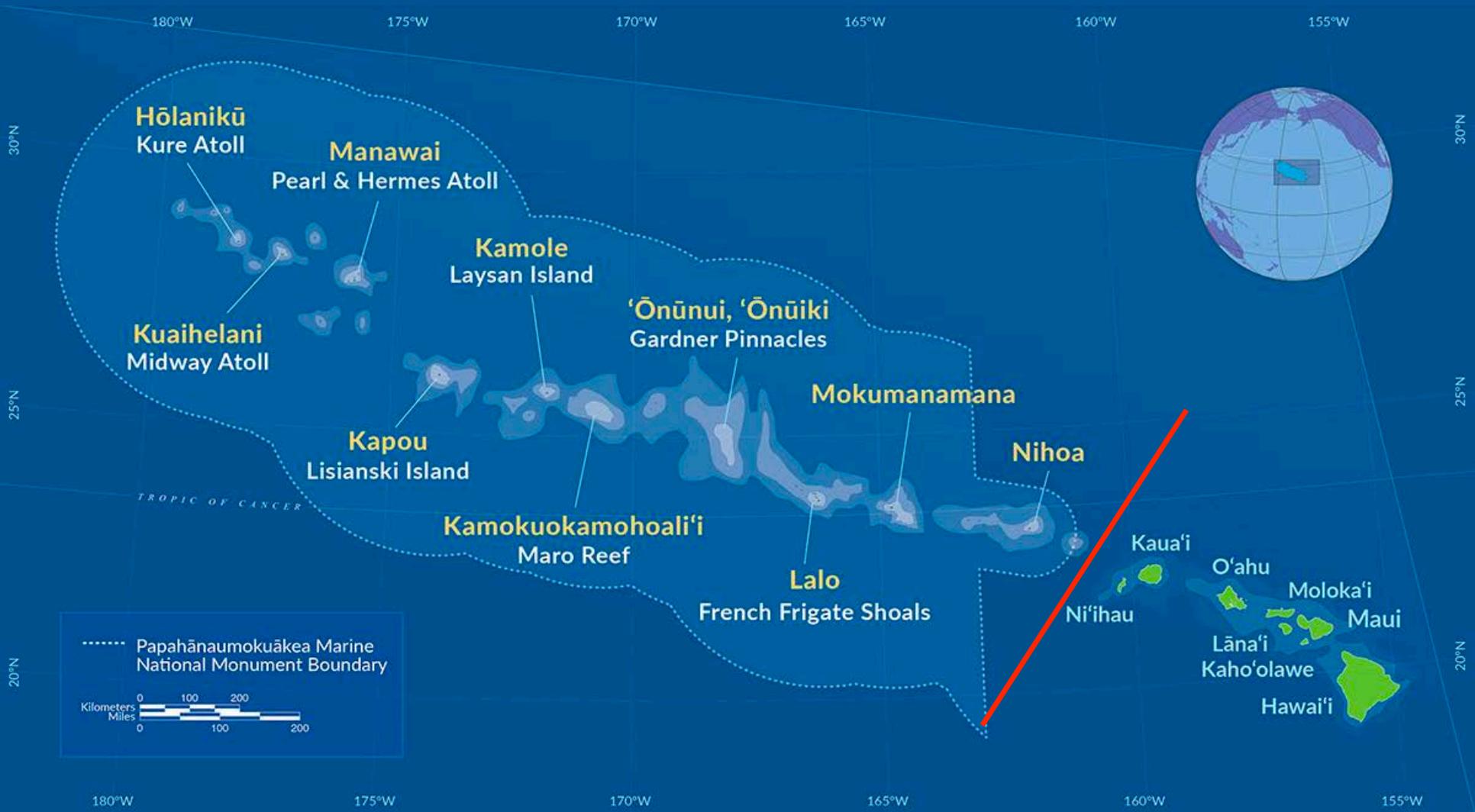
Hawaiian Archipelago



Hawaiian Archipelago



Hawaiian Archipelago



Archipelago

Island



Archipelago



Island



Genetic methods



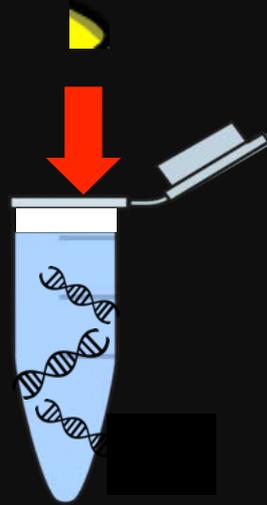
Collect tissue sample



Genetic methods



Collect tissue sample



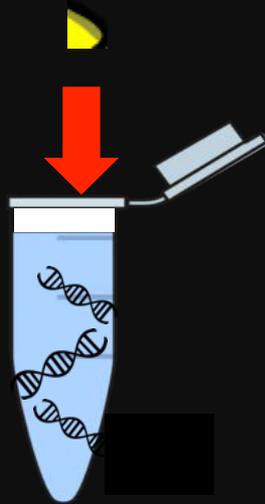
Extract & Isolate DNA



Genetic methods



Collect tissue sample



Extract & Isolate DNA



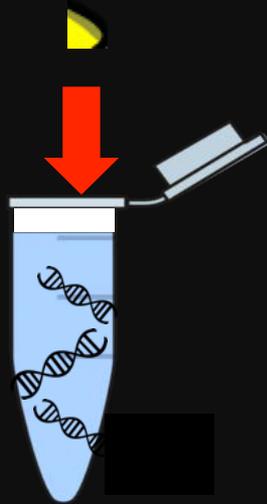
Sequence DNA



Genetic methods



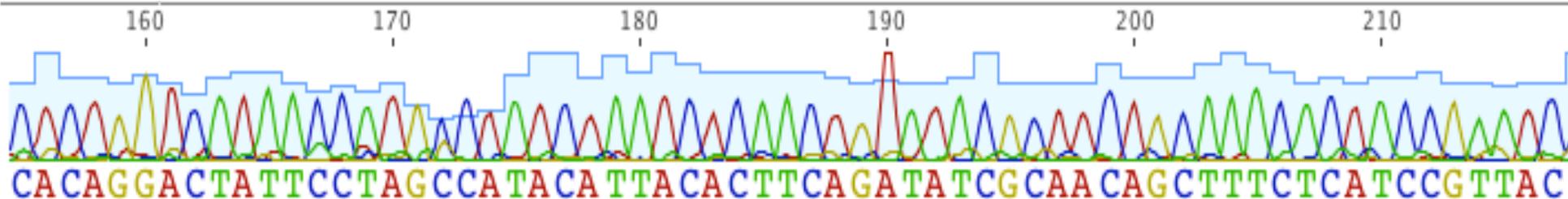
Collect tissue sample



Extract & Isolate DNA

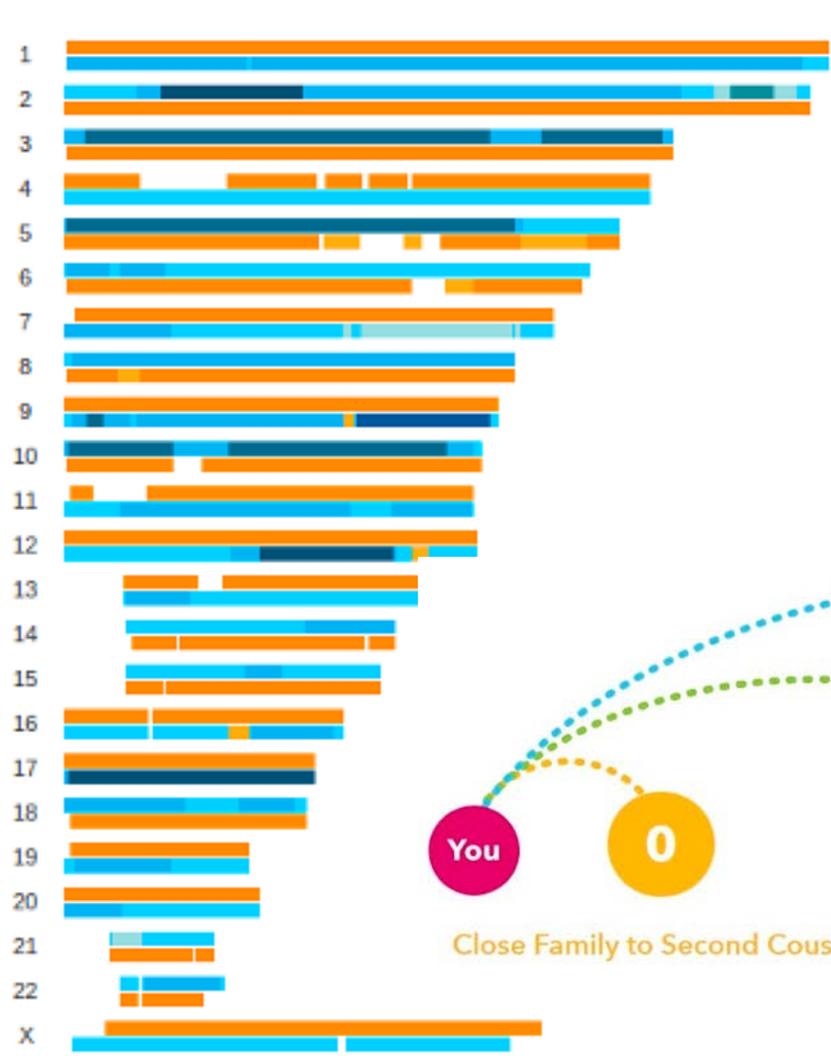


Sequence DNA



Using genetic methods

Chromosome View ▾ - Sub-regional Resolution +



Using genetic methods



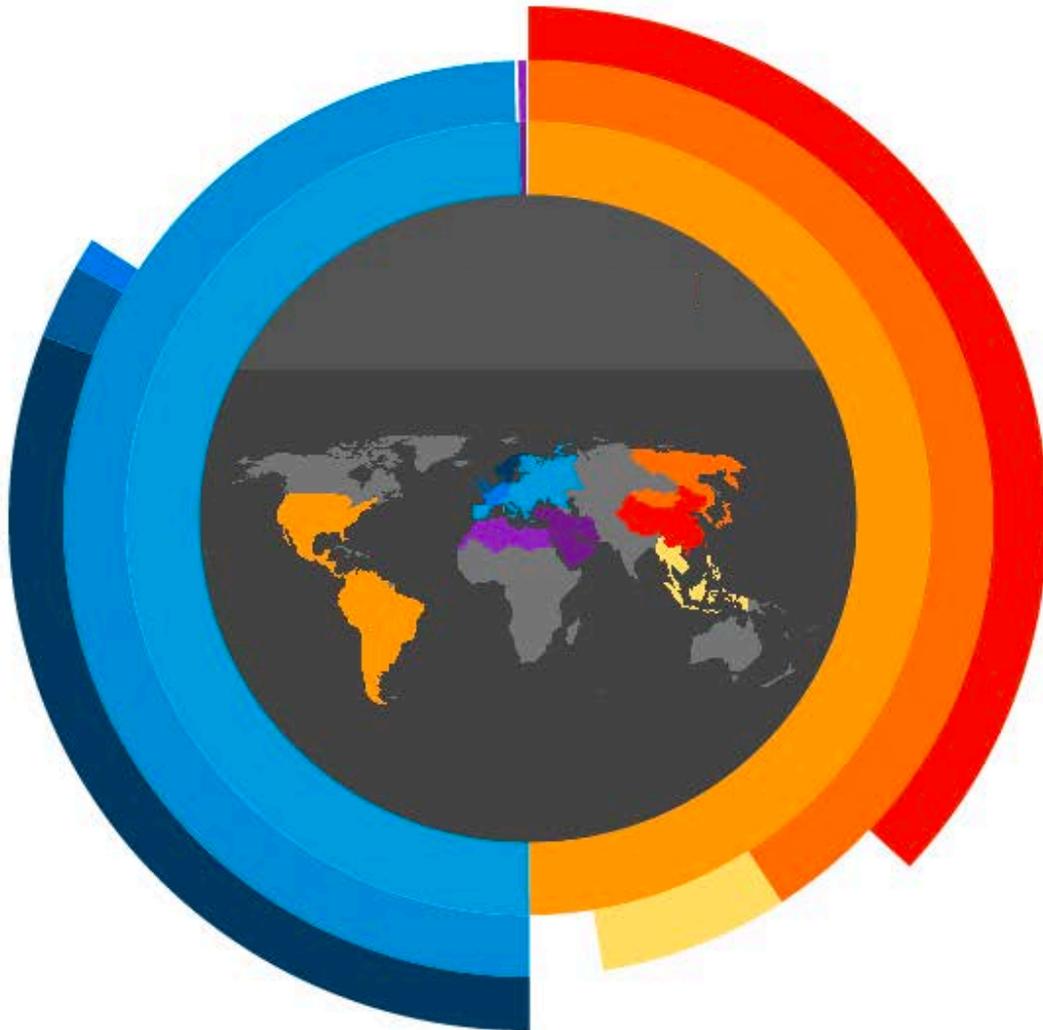
Map View



Sub-regional Resolution



Ancestry Composition tells you what percent of your DNA comes from each of 31 populations worldwide. This analysis includes DNA you received from all of your recent ancestors, on both sides of your family. The results reflect where your ancestors lived before the widespread migrations of the past few hundred years.



- 49.9% East Asian & Native American
 - East Asian
 - 36.9% Chinese
 - 3.9% Broadly East Asian
 - 6.6% Southeast Asian
 - 2.5% Broadly East Asian & Native American

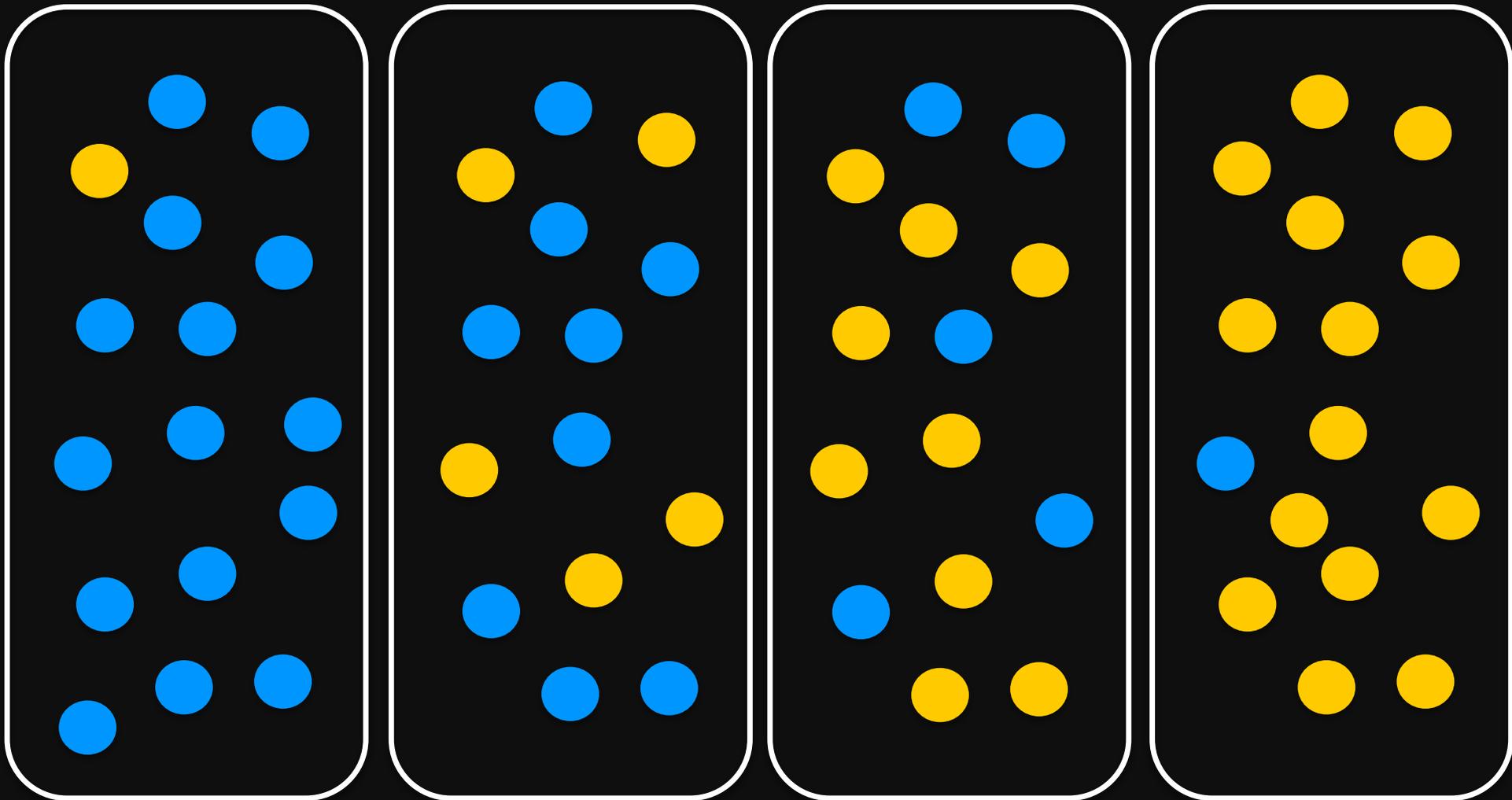
- 49.7% European
 - Northwestern European
 - 30.9% Scandinavian
 - 2.4% British & Irish
 - 1.0% French & German
 - 15.4% Broadly Northwestern European
 - 0.1% Broadly European

- 0.3% Middle Eastern & North African
- 0.3% North African

0.1% Unassigned

[show all populations](#)

Population Genetics Example



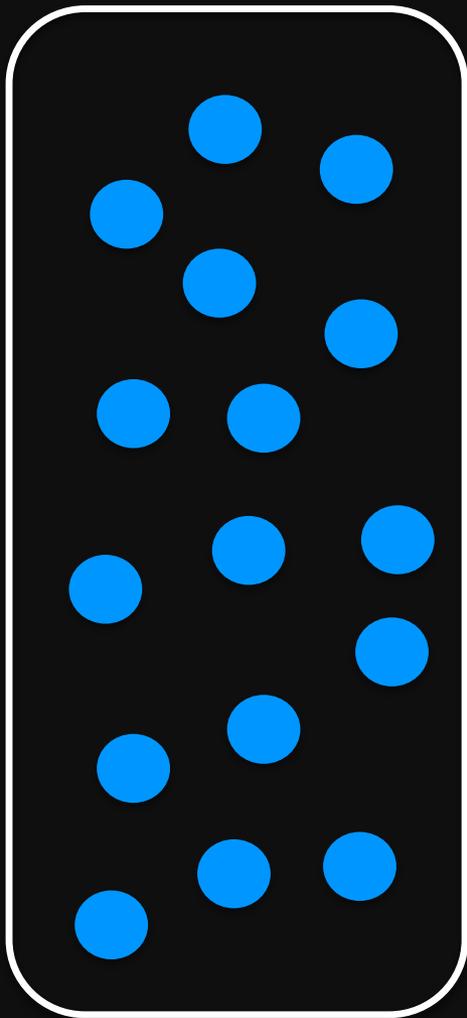
Pop 1

Pop 2

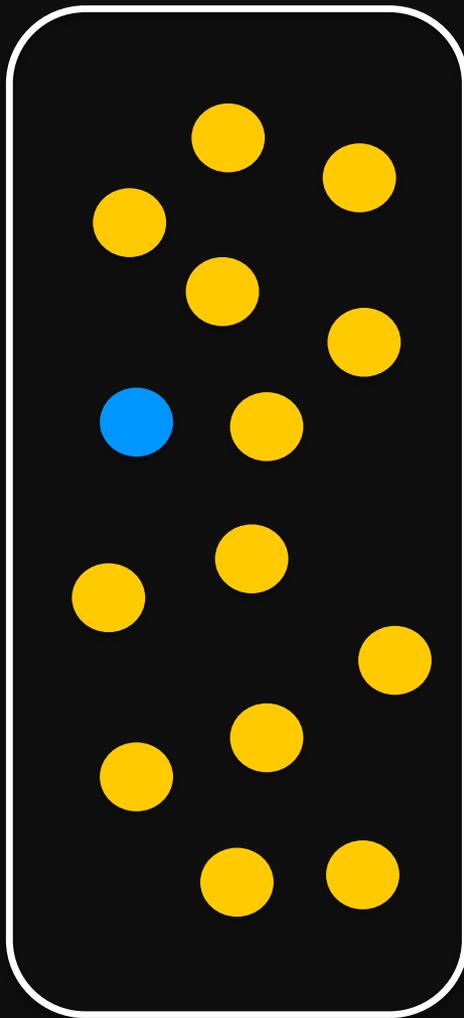
Pop 3

Pop 4

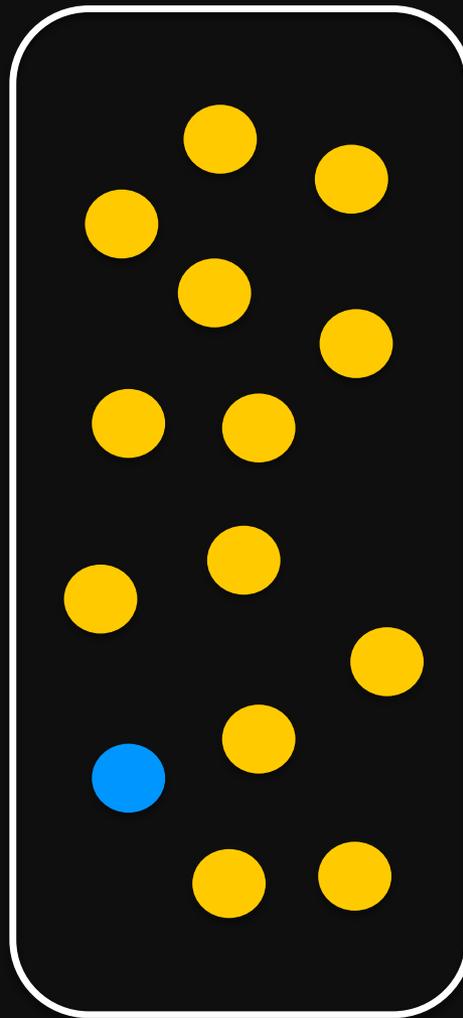
Population Genetics Example 2



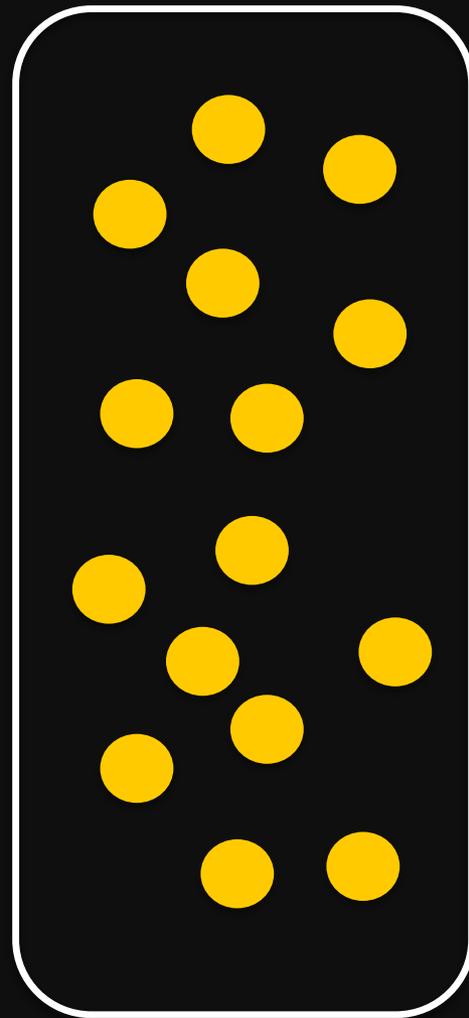
Pop 1



Pop 2

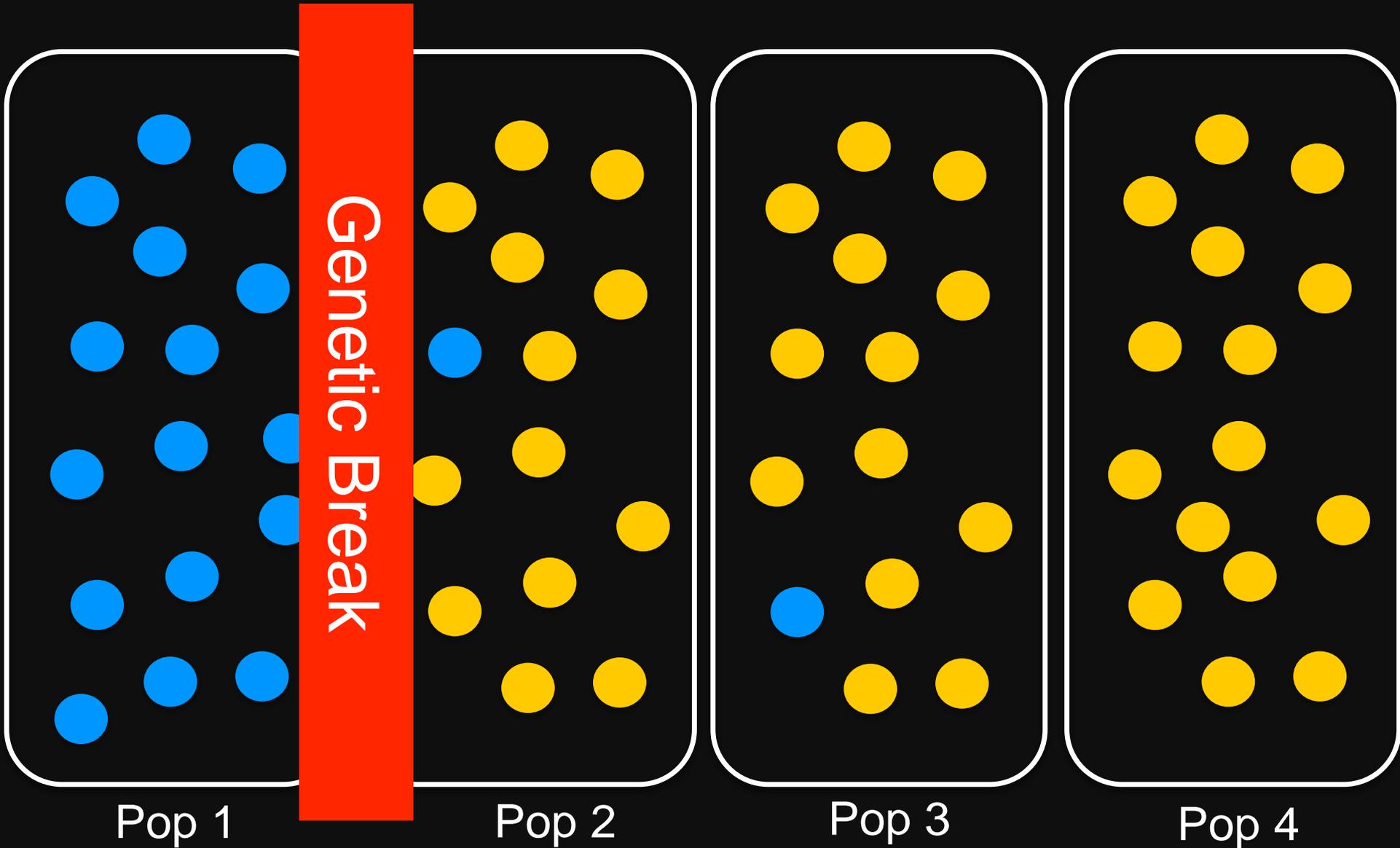


Pop 3

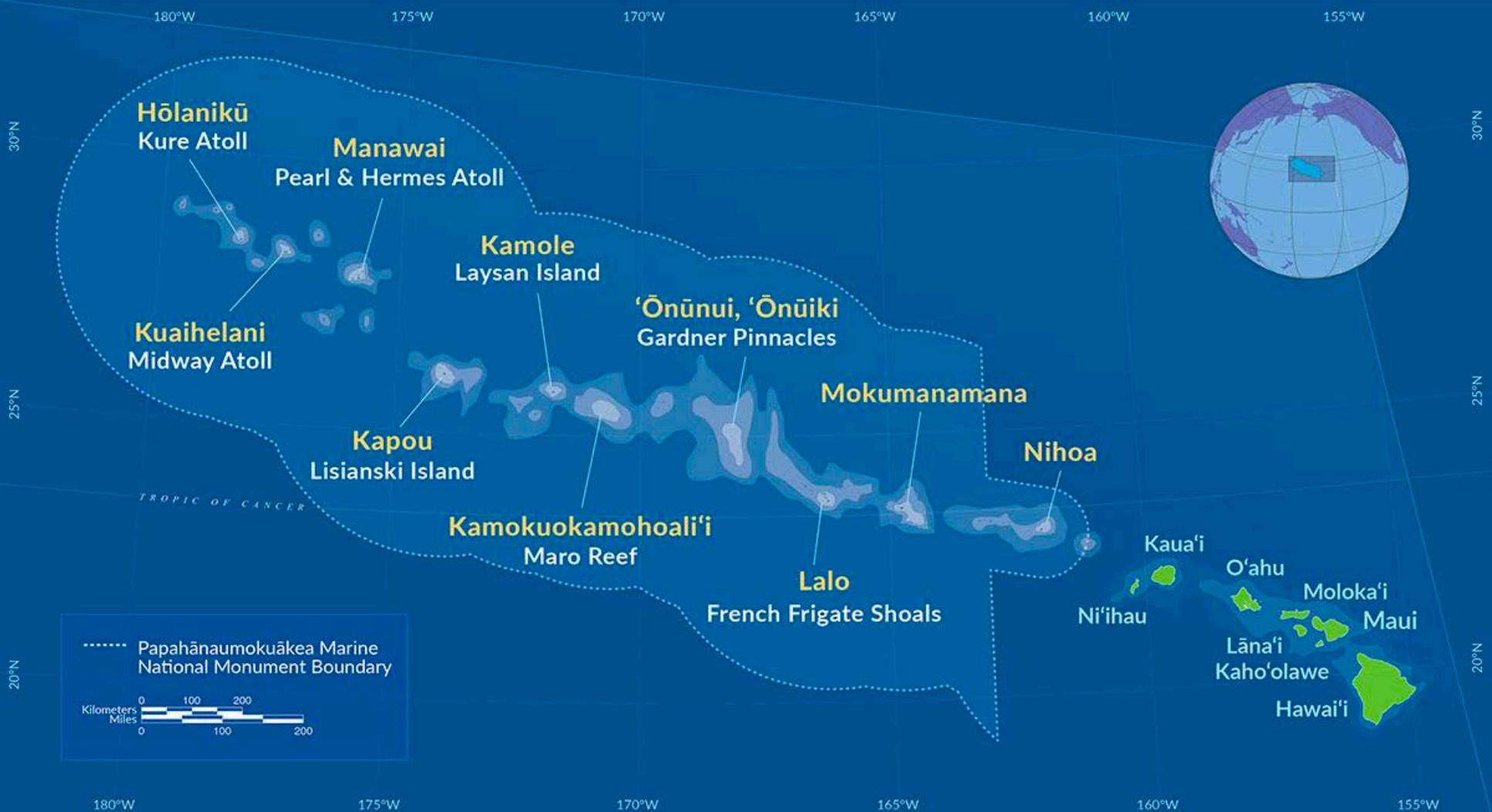


Pop 4

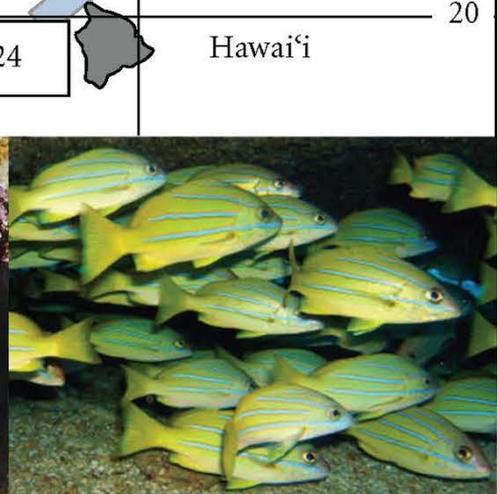
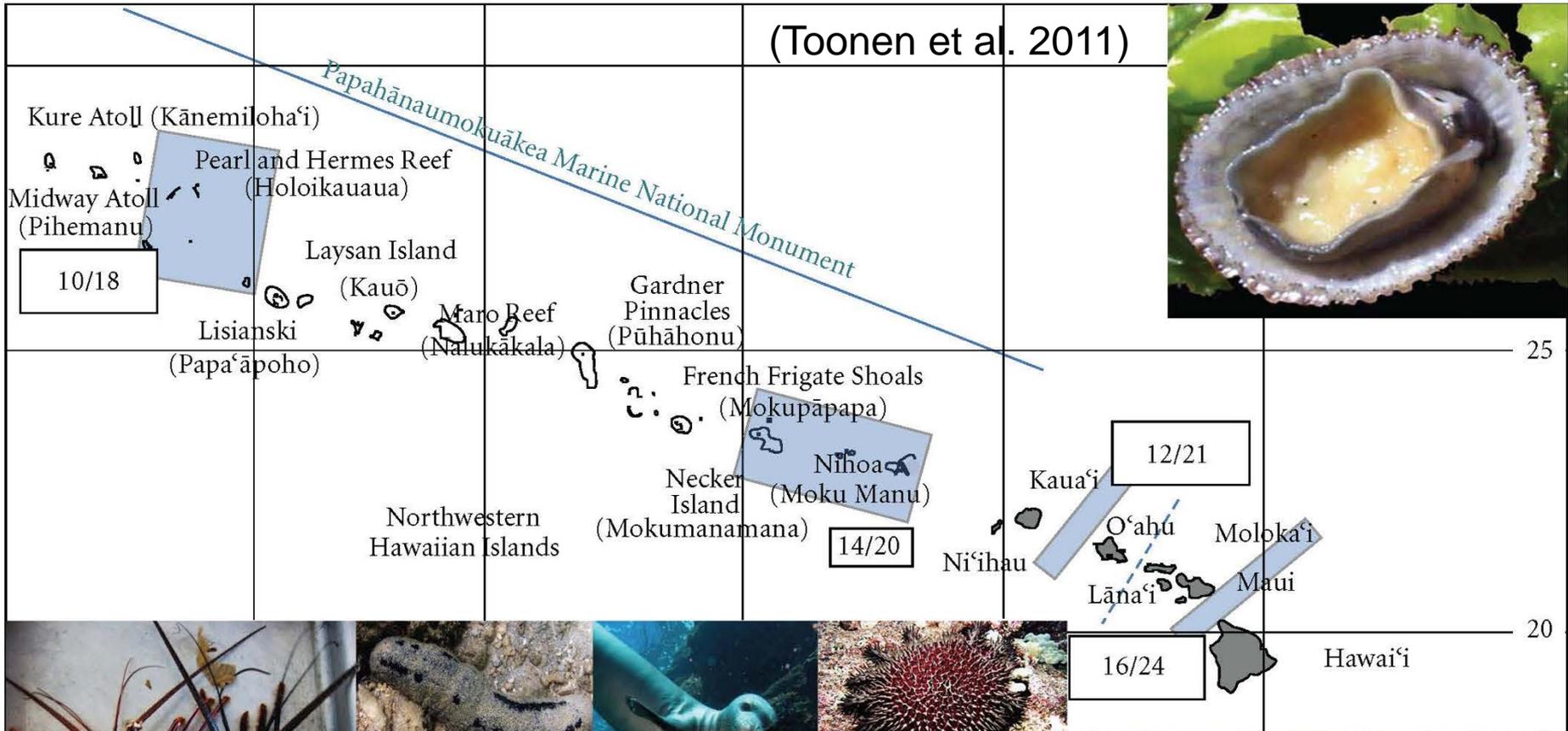
Population Genetics Example 2



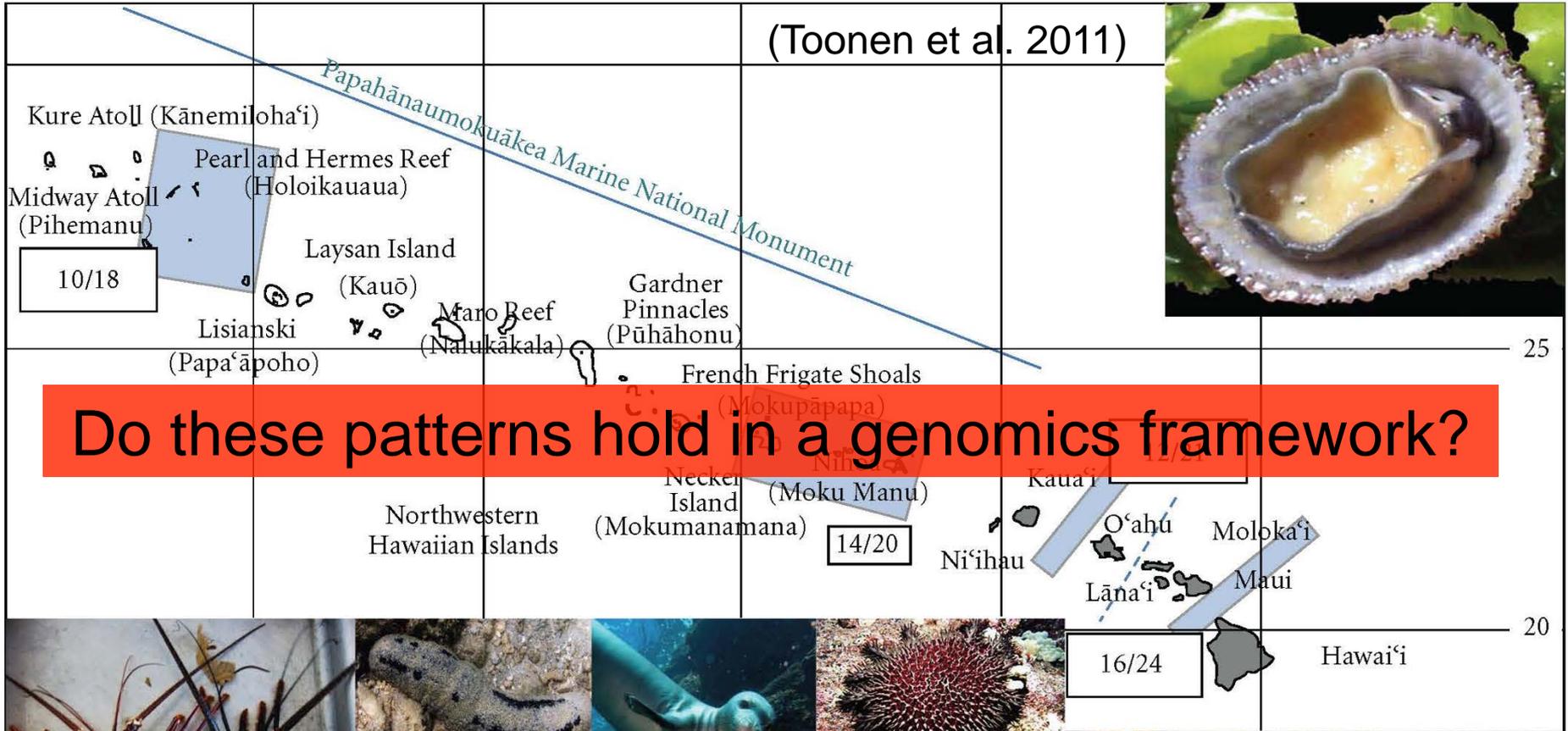
Hawaiian Archipelago



(Toonen et al. 2011)



(Toonen et al. 2011)



Manini (*Acanthurus triostegus*)



- PLD: 54-70 days
- Indo-Pacific

Kole (*Ctenocheatus strigosus*)



- PLD: 50-60 days
- Hawaiian Endemic

Manini (*Acanthurus triostegus*)



- PLD: 54-70 days
- Indo-Pacific = high dispersal

Kole (*Ctenocheatus strigosus*)



- PLD: 50-60 days
- Hawaiian Endemic = low dispersal

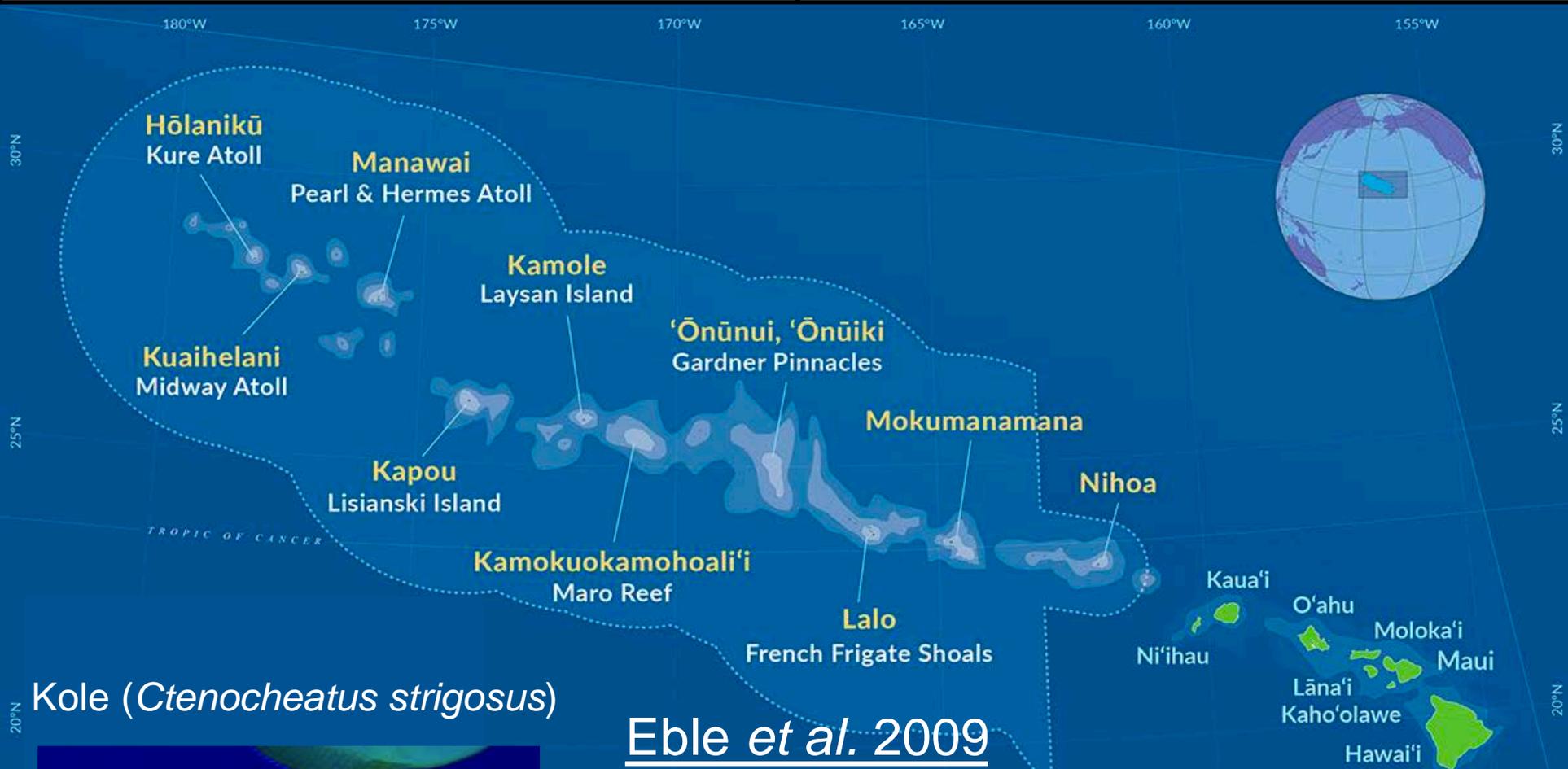
Manini (*Acanthurus triostegus*)

- PLD: 54-70 days
- Indo-Pacific = high dispersal

Kole (*Ctenocheatus strigosus*)

- PLD: 50-60 days
- Hawaiian Endemic = low dispersal

Do these species show similar connectivity patterns across the Hawaiian Archipelago?

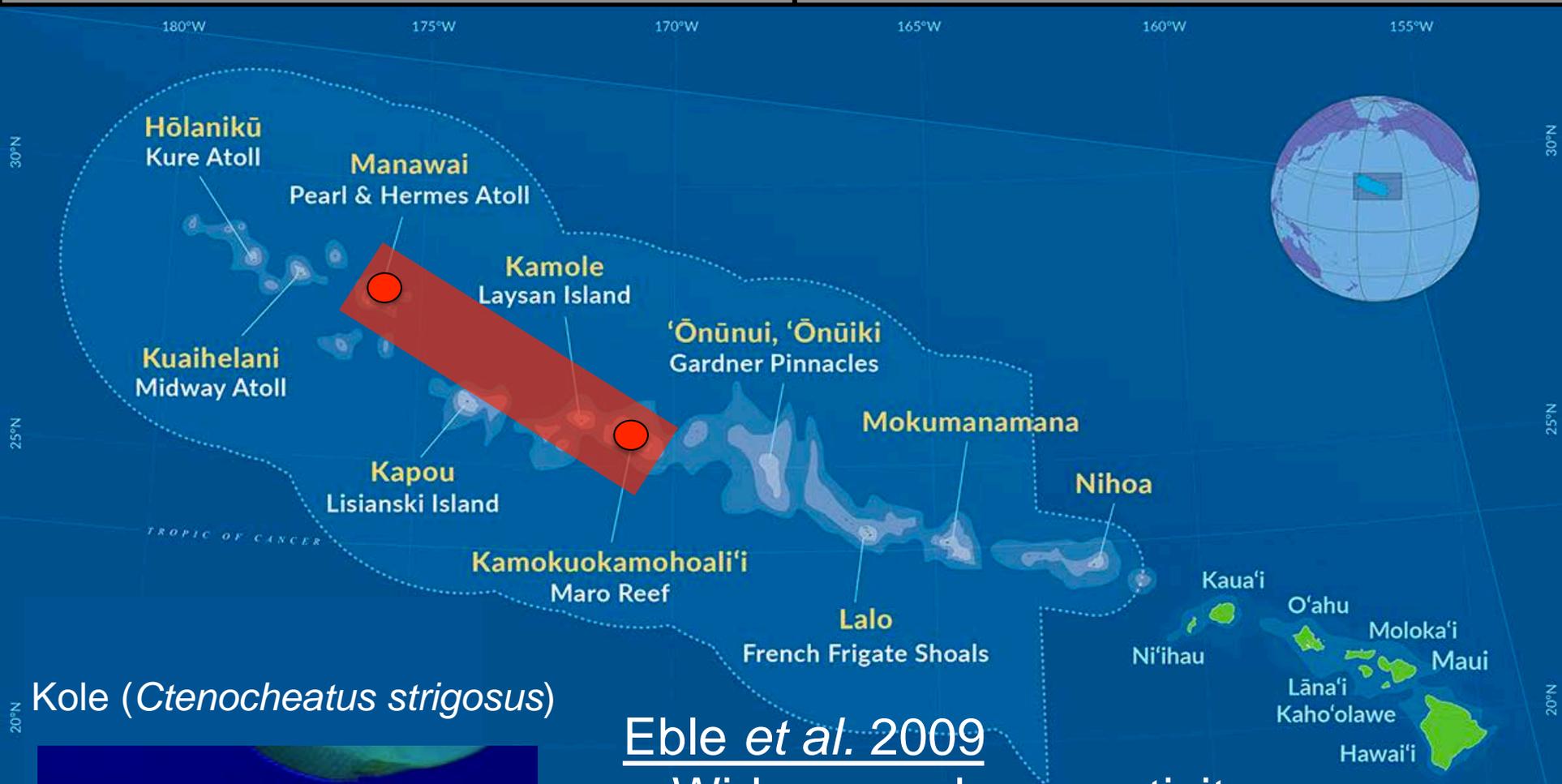


Kōle (*Ctenocheatus strigosus*)



Eble et al. 2009

- One mitochondrial DNA marker



Kole (*Ctenocheatus strigosus*)

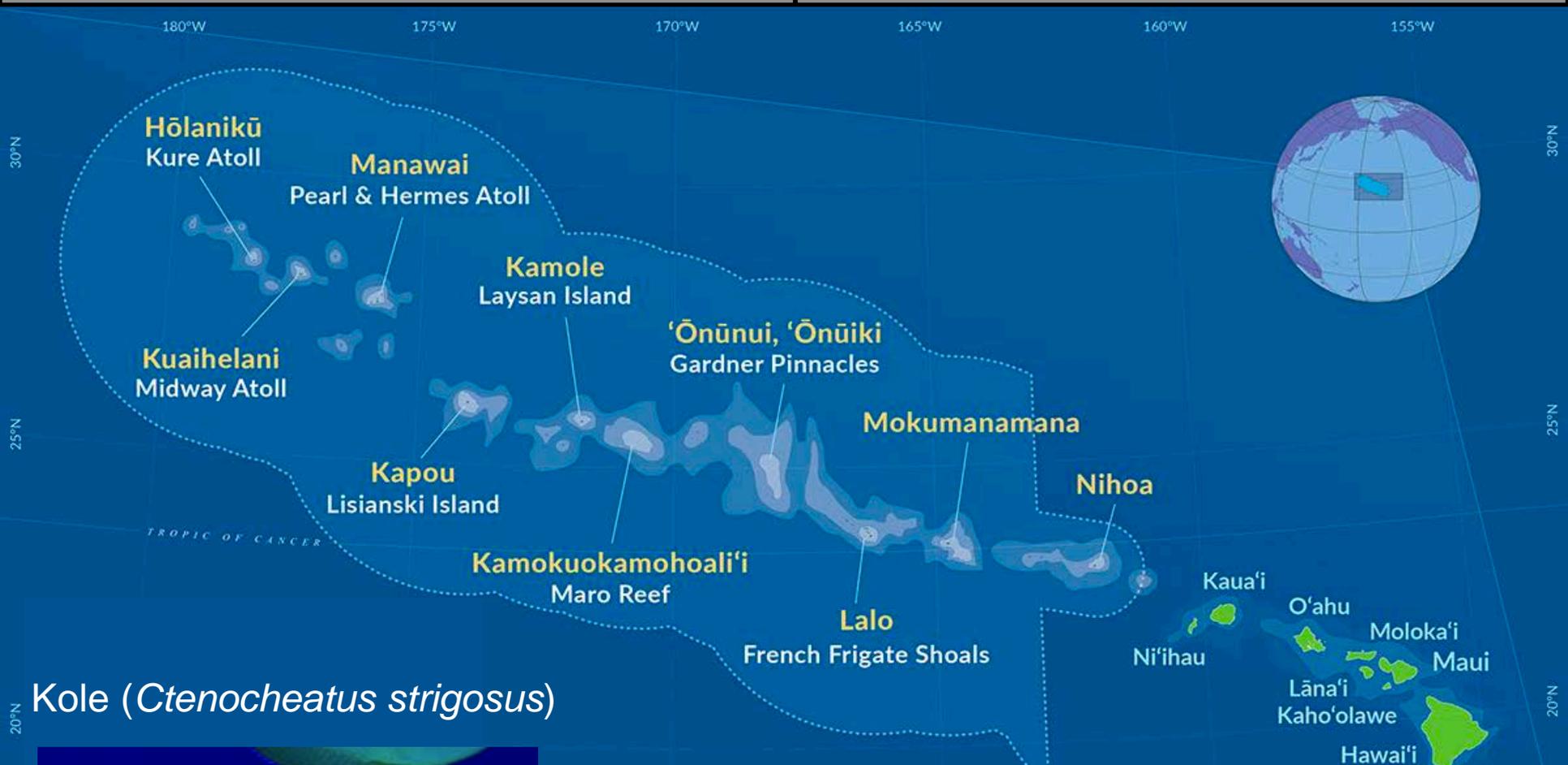


Eble et al. 2009

- Wide spread connectivity
- Pearl & Hermes, Maro Reef isolated from the rest of the archipelago

Archipelago

Island

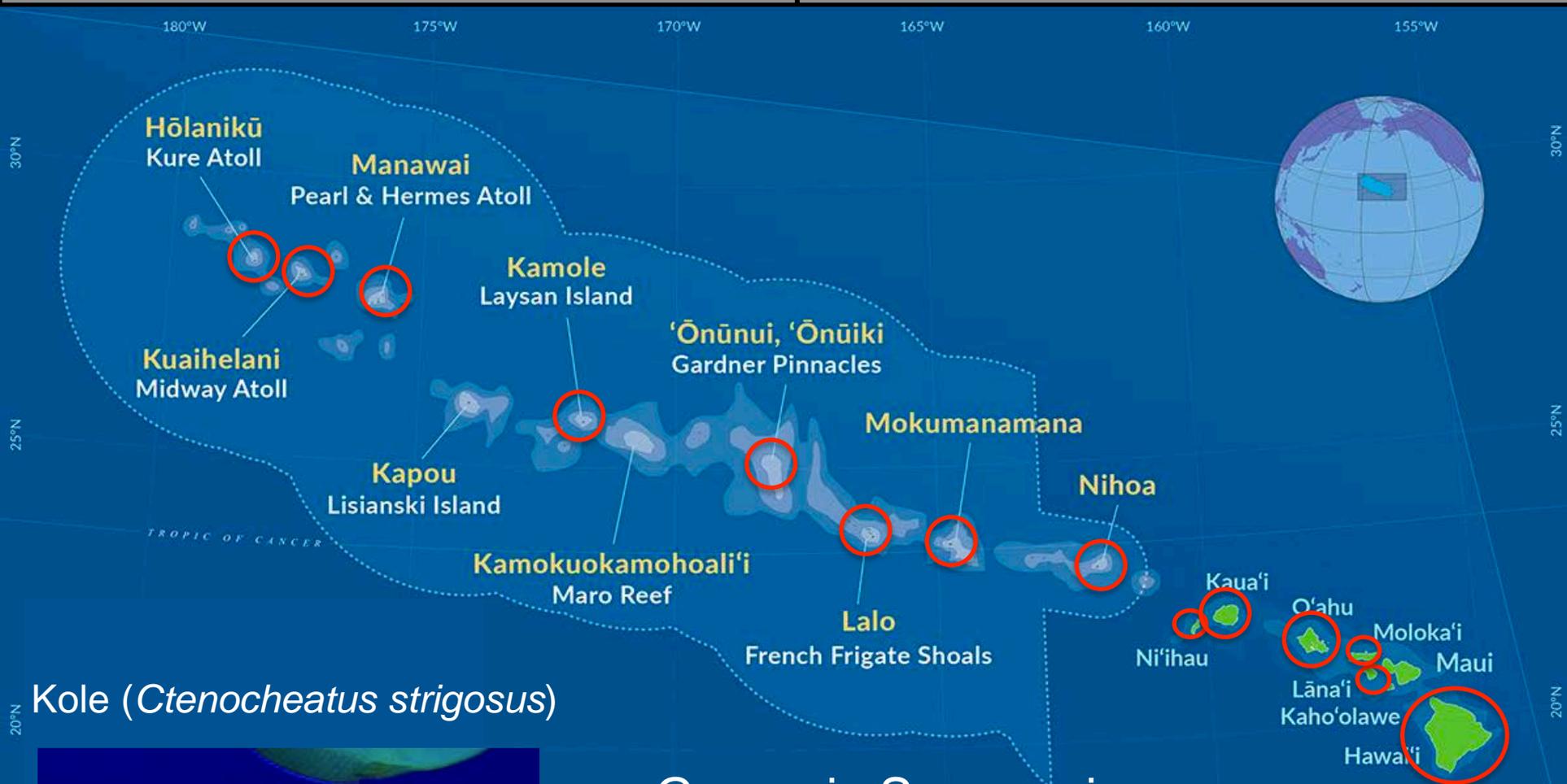


Kole (*Ctenocheatus strigosus*)



© Keoki Stender

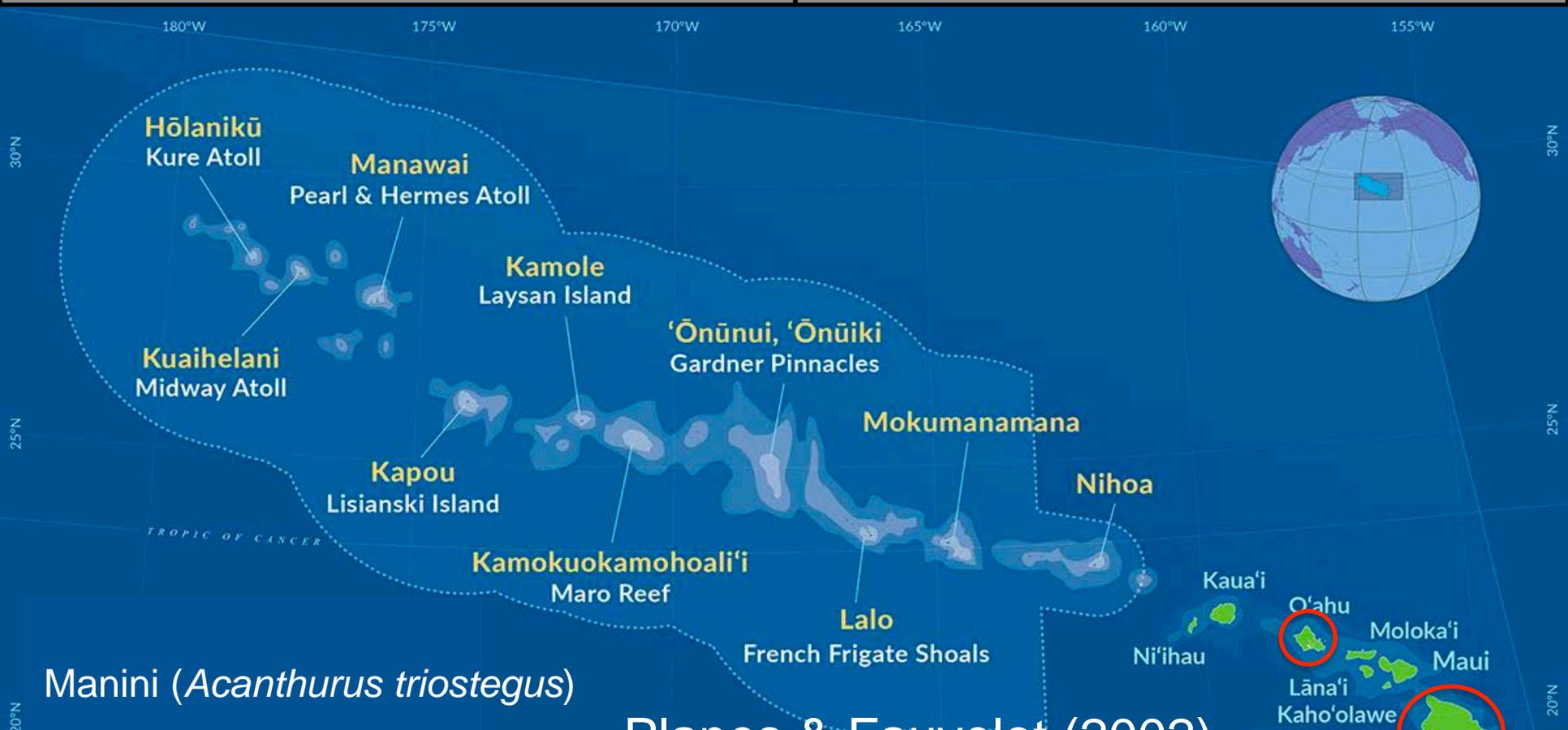
- Genomic Sequencing



Kole (*Ctenocheatus strigosus*)



- Genomic Sequencing
- Each Island is genetically distinct
- Each island needs to be managed independently

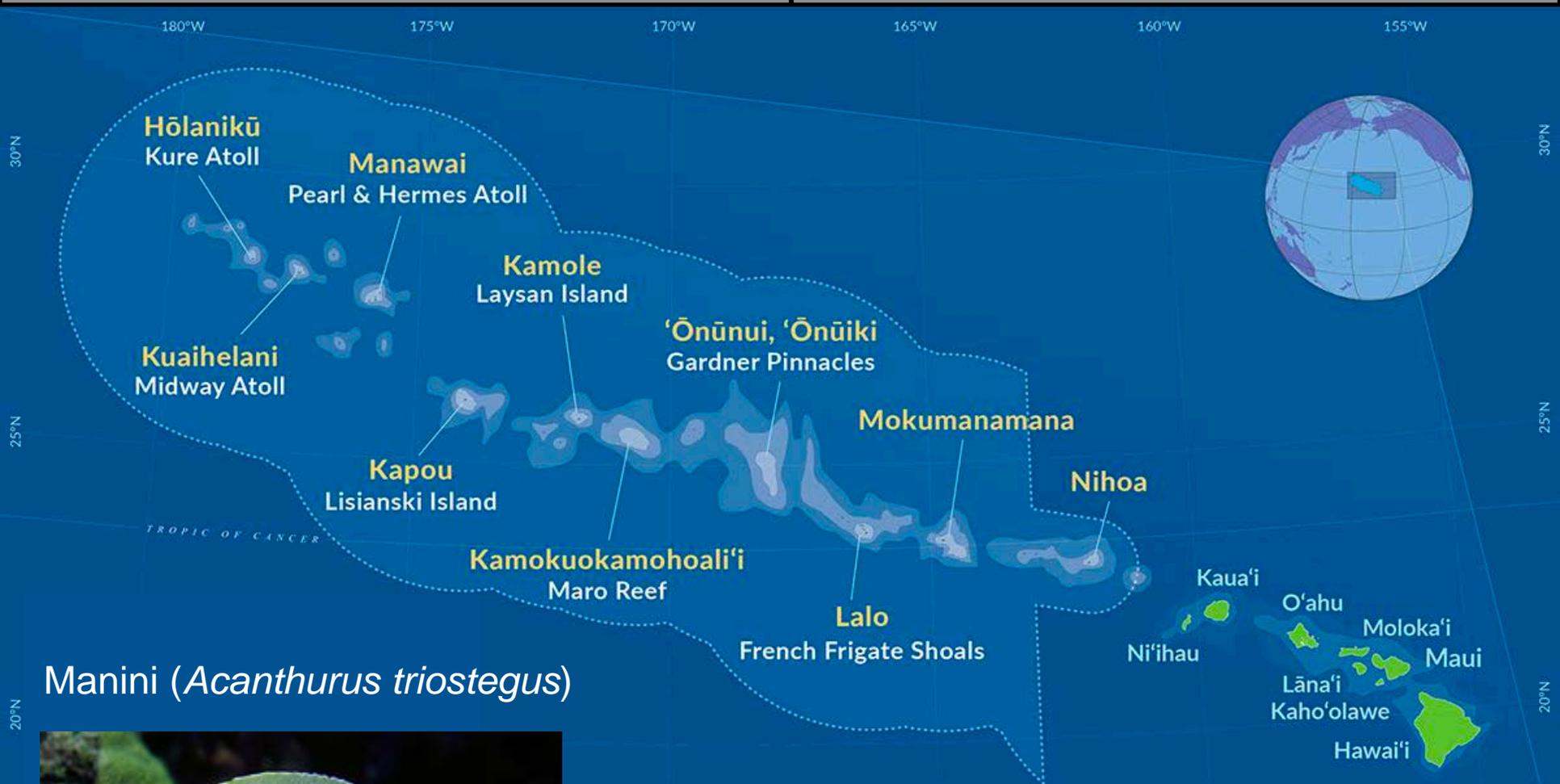


Manini (*Acanthurus triostegus*)



Planes & Fauvelot (2002)

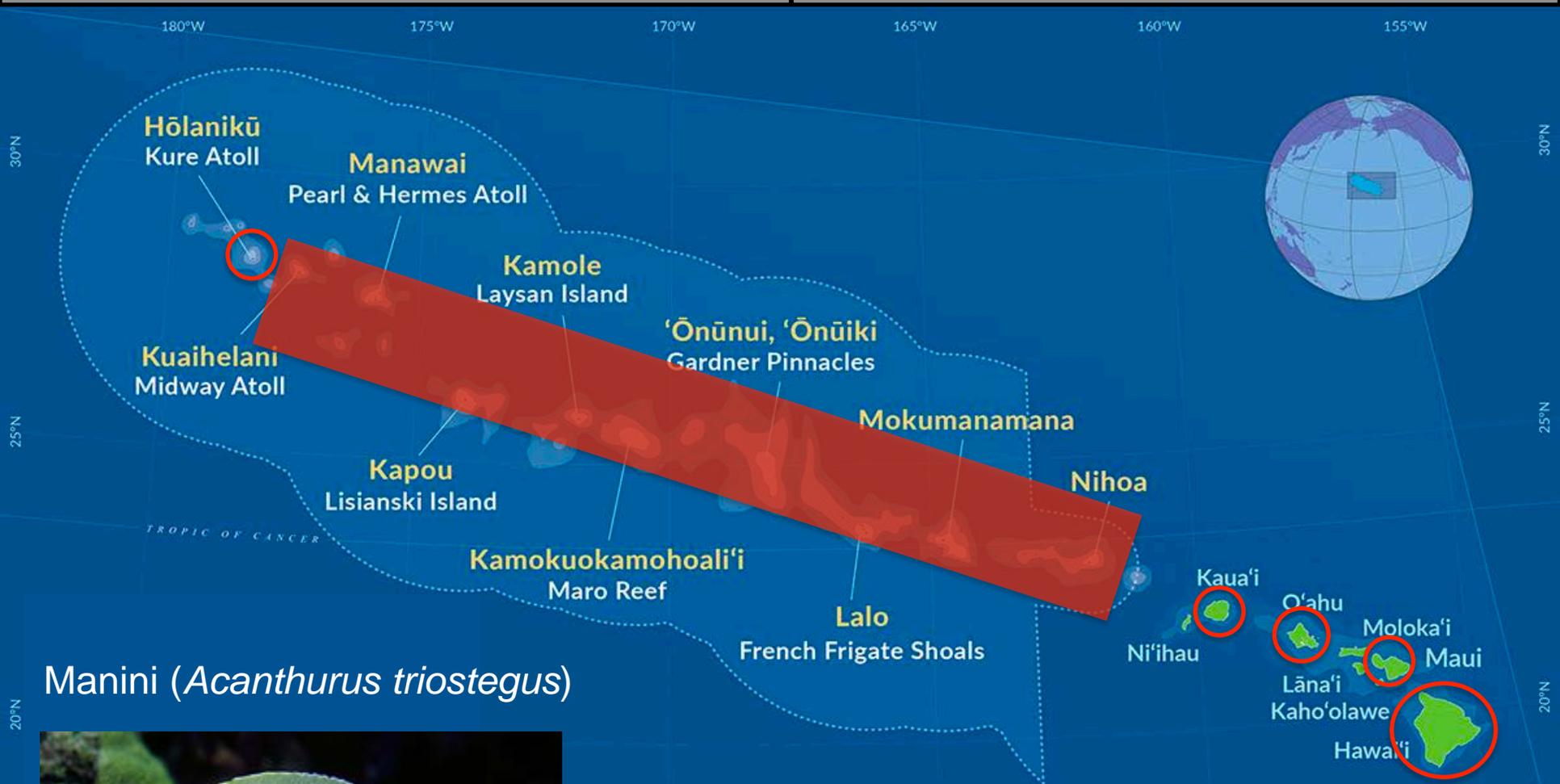
- O'ahu and Hawai'i Island genetically distinct



Manini (*Acanthurus triostegus*)



- Genomic Sequencing



Manini (*Acanthurus triostegus*)



- Genomic Sequencing
- Each MHI, Kure; isolated populations
- NWHI + Johnston genetically homogenous

Conclusions: Archipelago

Manini (*Acanthurus triostegus*)



Kole (*Ctenocheatus strigosus*)



Conclusions: Archipelago

Manini (*Acanthurus triostegus*)



Kole (*Ctenocheatus strigosus*)



Genomics provides finer scale resolution for identifying connectivity patterns

Conclusions: Archipelago

Manini (*Acanthurus triostegus*)



Kole (*Ctenocheatus strigosus*)



Genomics provides finer scale resolution for identifying connectivity patterns

- Island-by-island isolation
 - First account in Hawaiian fishes

Conclusions: Archipelago

Manini (*Acanthurus triostegus*)



Kole (*Ctenocheatus strigosus*)



Genomics provides finer scale resolution for identifying connectivity patterns

- Widespread connectivity in NWHI
- Island-by-island isolation
 - First account in Hawaiian fishes

Conclusions: Archipelago

Manini (*Acanthurus triostegus*)



Kole (*Ctenocheatus strigosus*)

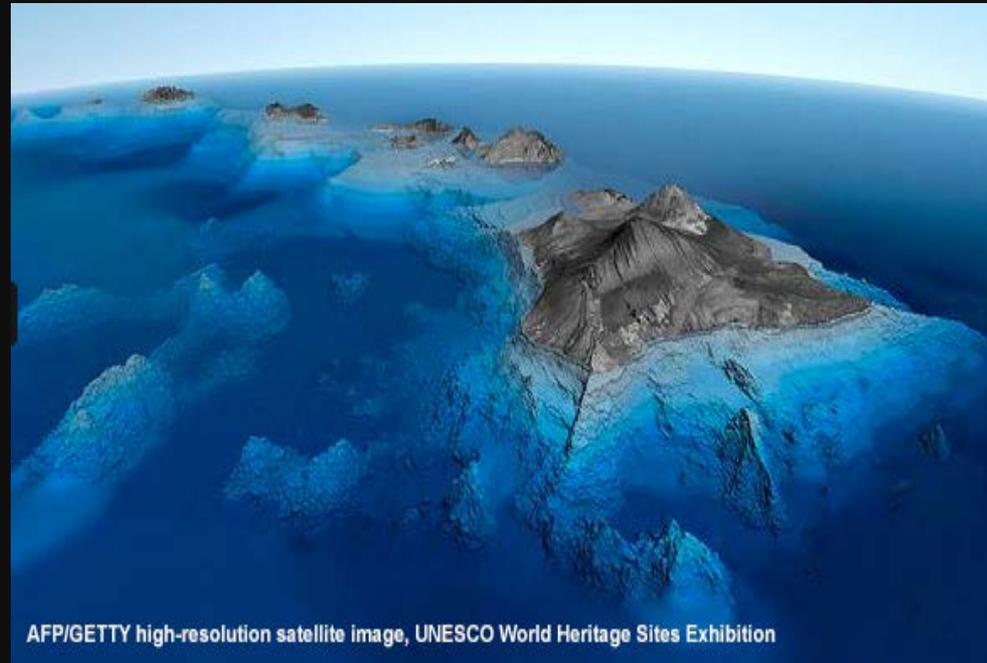


Management Implications

- MHI are separate populations from the NWHI

Smaller spatial scales: Island

- Archipelago assessment shows that each MHI is distinct
- Island assessment requires finer scale resolution
 - Parentage analysis



Parentage Analysis



Recreational Fisheries in Hawai'i

- Heavy pressure of fish communities
- Not well regulated
- Estimated that 31% of Hawai'i residents participate in recreational fishing (Hamnett et al. 2006)
- 36% of the total catch is attributed to recreational harvest (WestPac, 2002)



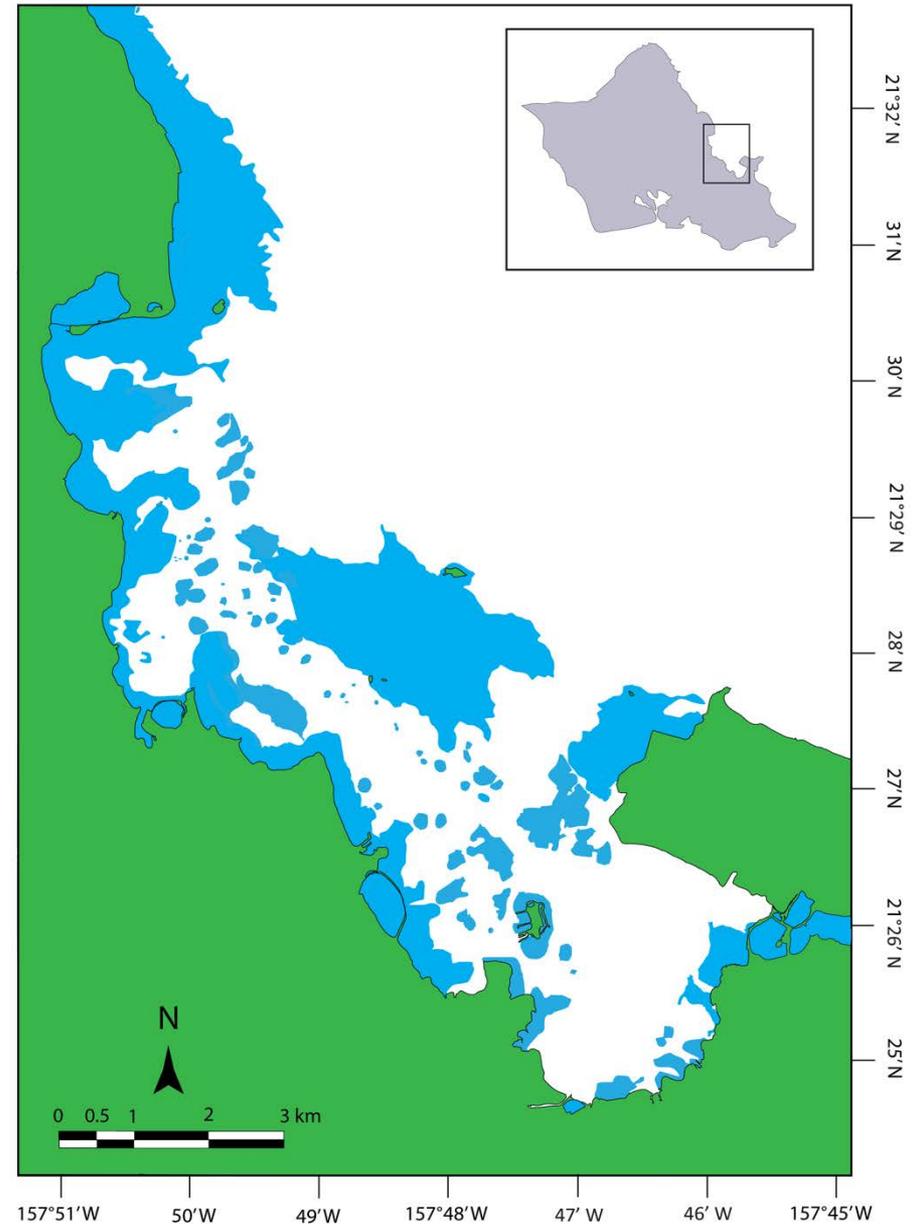
Problems associated with under-managed fisheries

- Lead to overfishing
- Prohibit long-term sustainability
 - Inability to maintain food security



Windward O'ahu

- Identify connectivity and dispersal patterns
- Initiated by Native Hawaiian community leaders to better understand connectivity of resources



Manini, *Acanthurus triostegus*

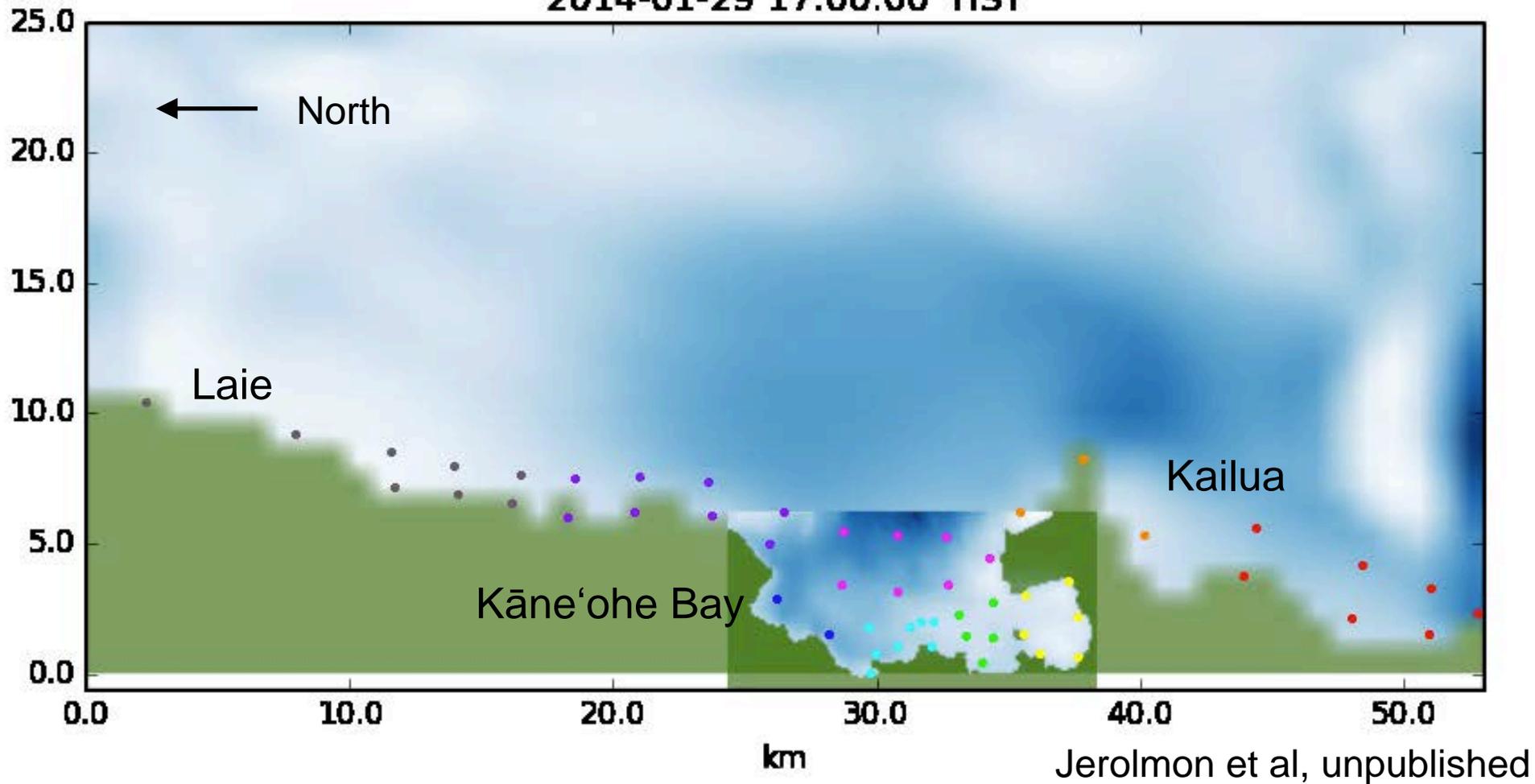


- Identified by communities leaders as an important fish stock
- Abundant throughout Hawai'i
- Heavily targeted by recreational fishers



Windward Coast of O'ahu

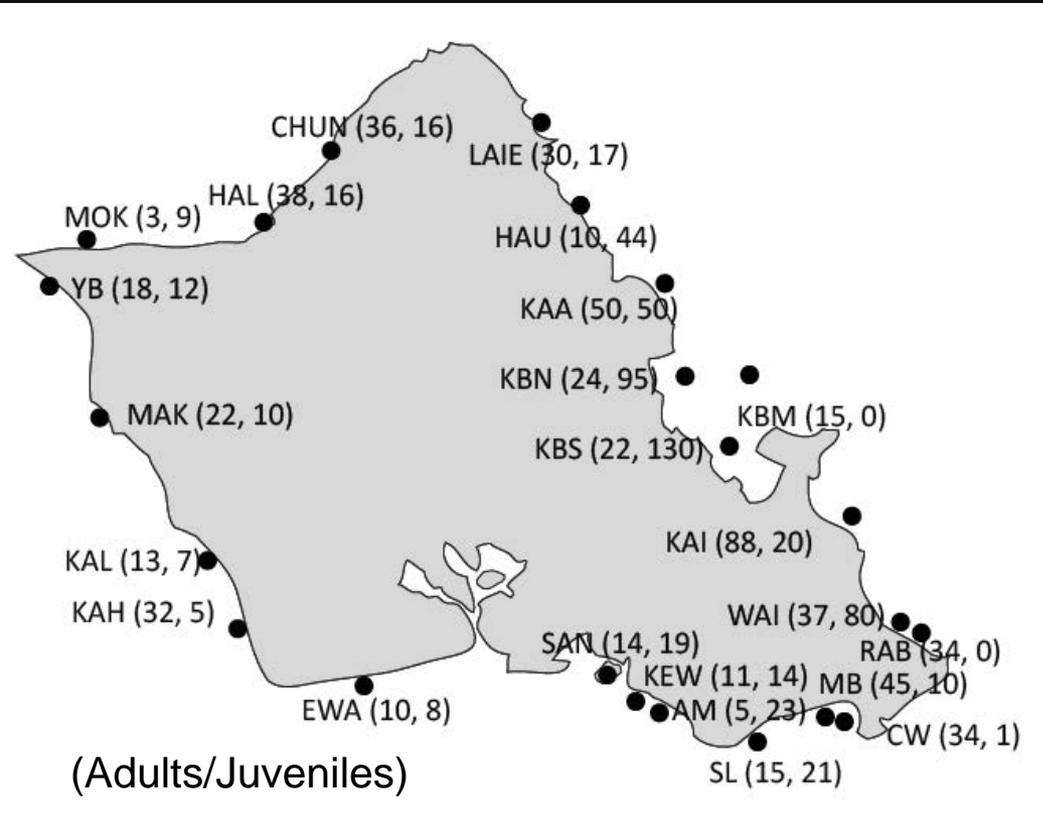
2014-01-29 17:00:00 HST



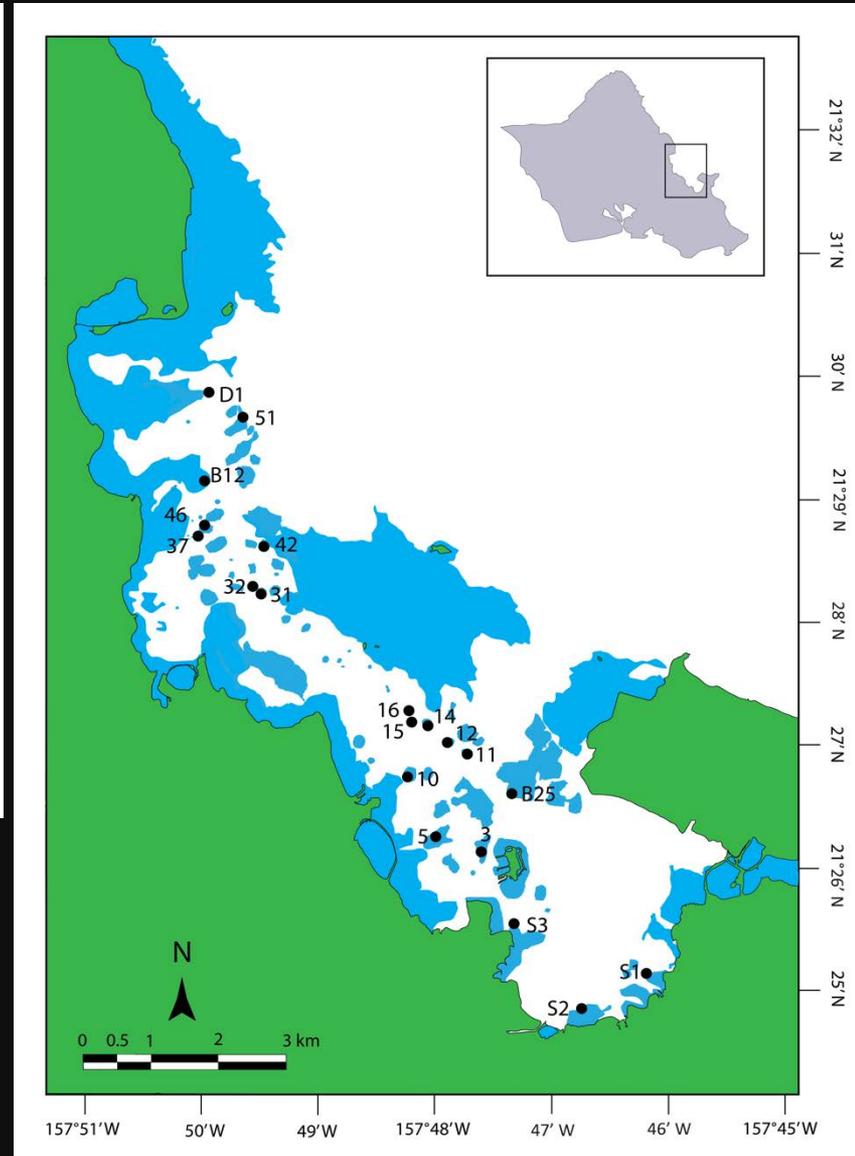
Goal

- Identify dispersal pathways
 - Source and sink populations
- Identify areas that may be vulnerable to fishing pressure

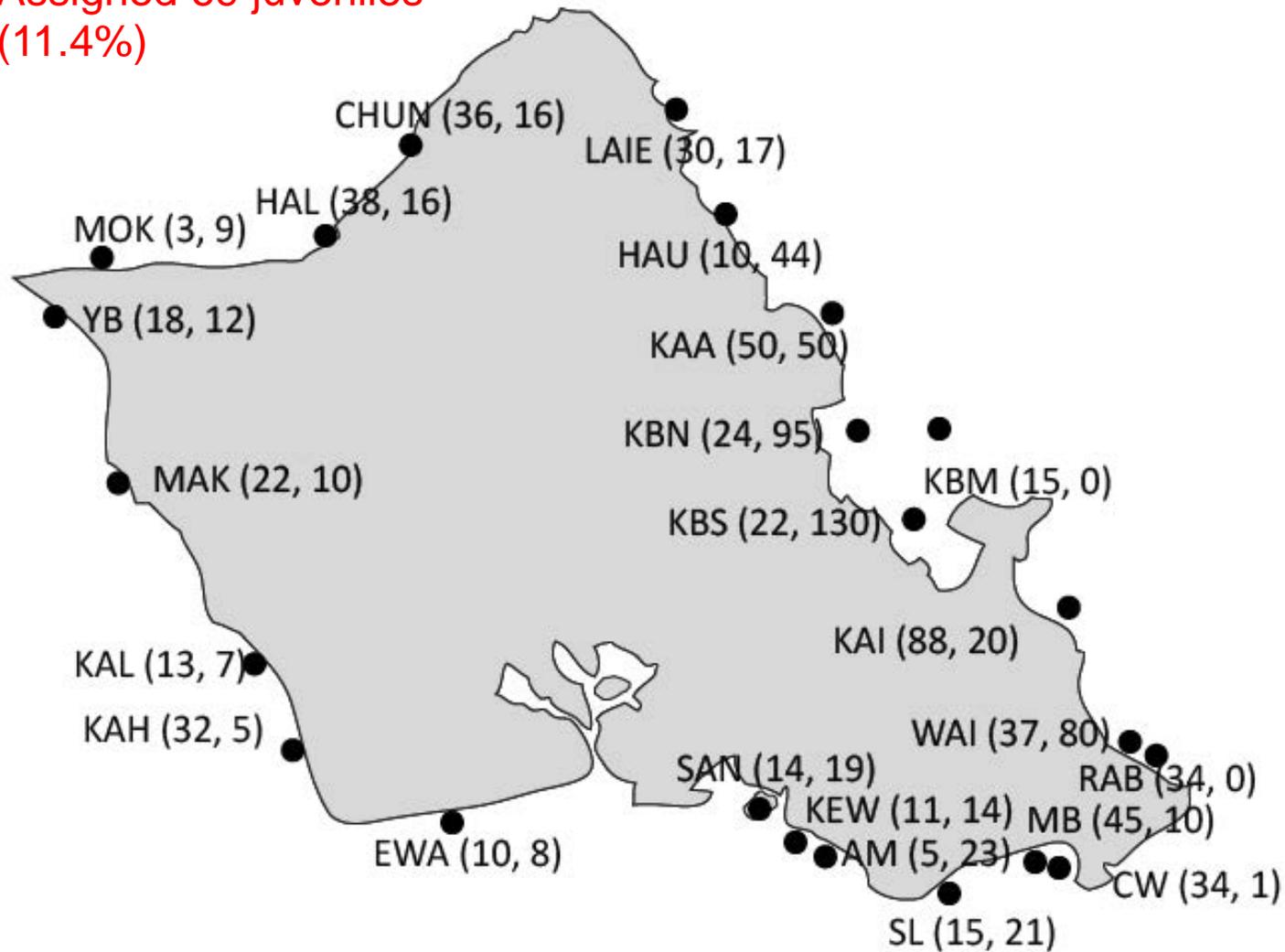
Collection Sites



- Collected 606 adults, 607 juveniles



Assigned 69 juveniles
(11.4%)



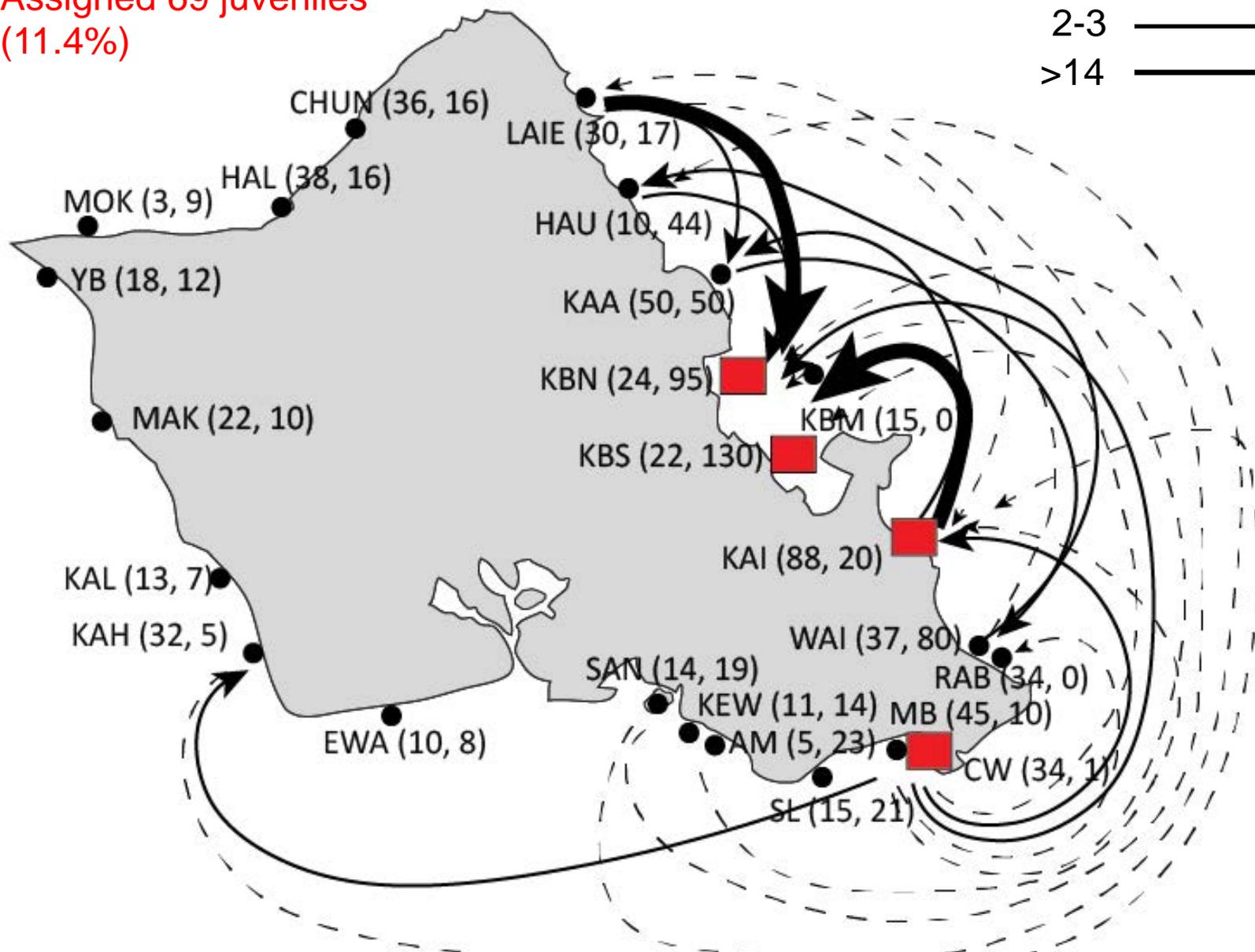
(Adults/Juveniles)

Archipelago

Island

Assigned 69 juveniles
(11.4%)

1
2-3
>14



(Adults/Juveniles)

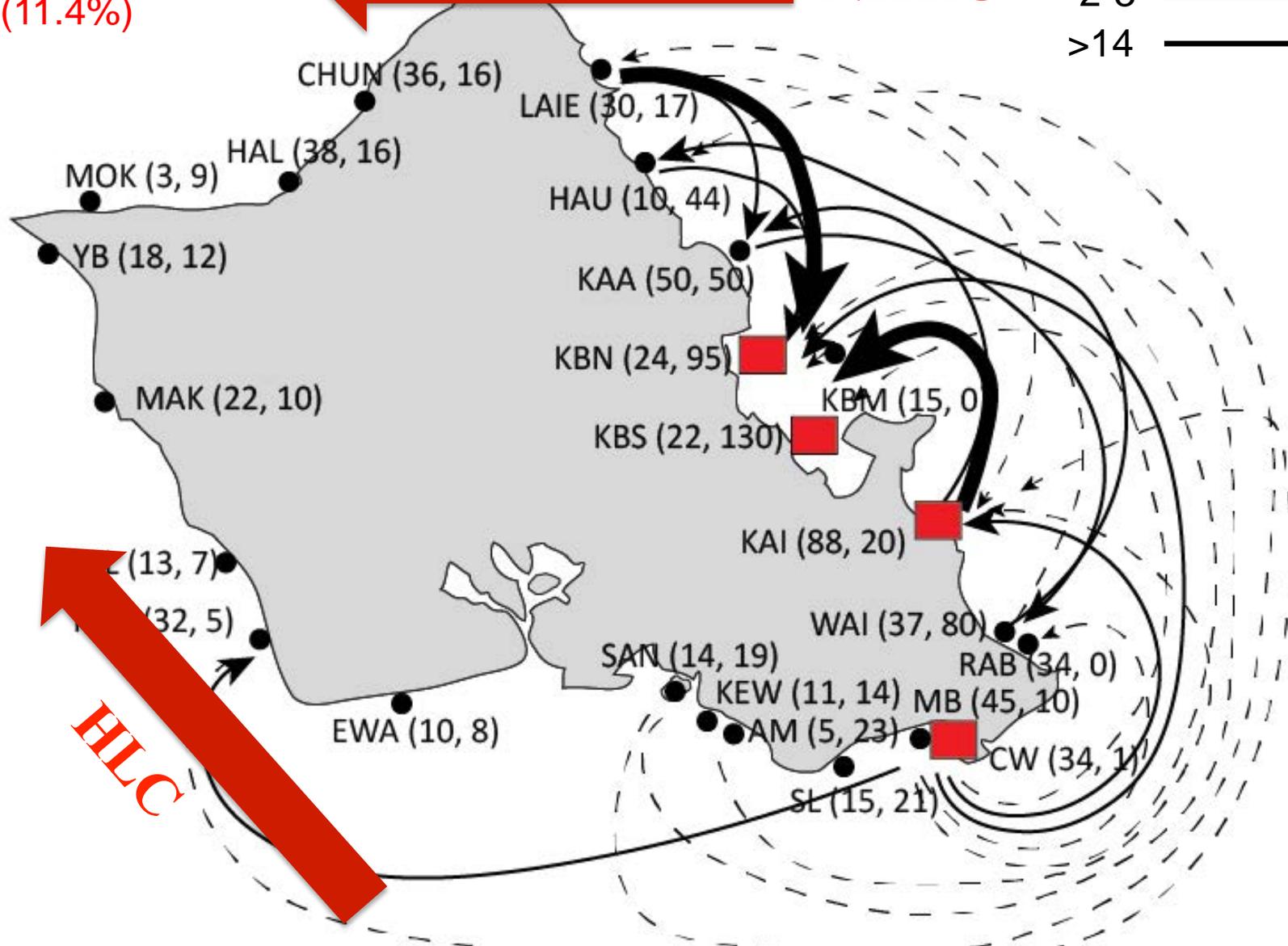
Archipelago

Island

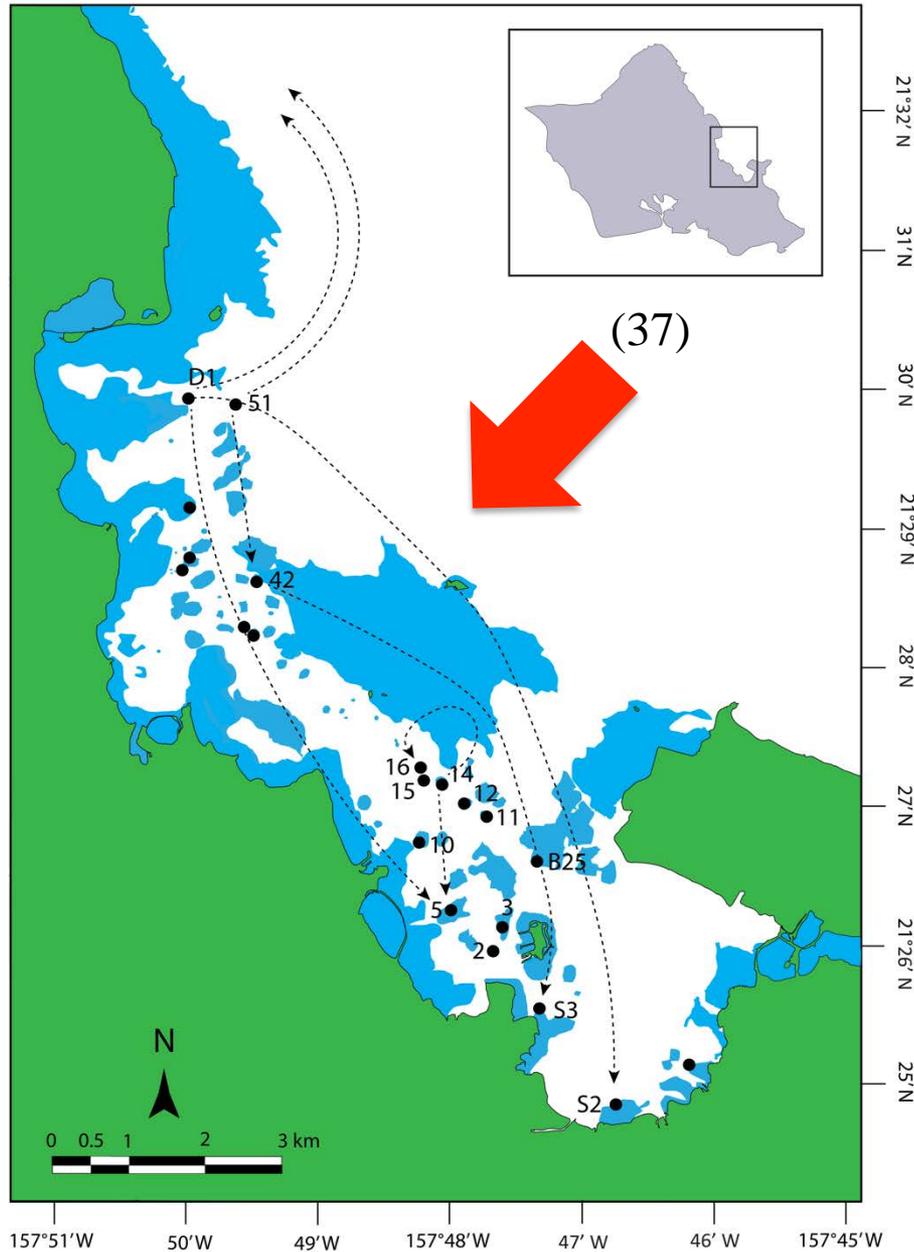
Assigned 69 juveniles
(11.4%)

NHRC

1	_____
2-3	=====
>14	=====



(Adults/Juveniles)



- Six occurrences of retention inside Kāneʻohe Bay
- Two instances of dispersal out of the bay (Hauʻula, Lāʻie)

Conclusions: Island

- Majority of larvae do not disperse far
 - Most dispersal is limited to the <30 km
 - Some indication of local retention
- East O‘ahu is a source for recruitment
 - Kāne‘ohe Bay is dependent on recruitment from outside of the bay
 - Management Implications
 - Not many adults within the bay

Spatial Scales



Archipelago
Hawaiian Archipelago

Island
O'ahu

Spatial Scales



Archipelago
Hawaiian Archipelago

Genomics provides finer scale resolution;
MHI are separate populations from NWHI

Island
O'ahu

Spatial Scales



Archipelago
Hawaiian Archipelago

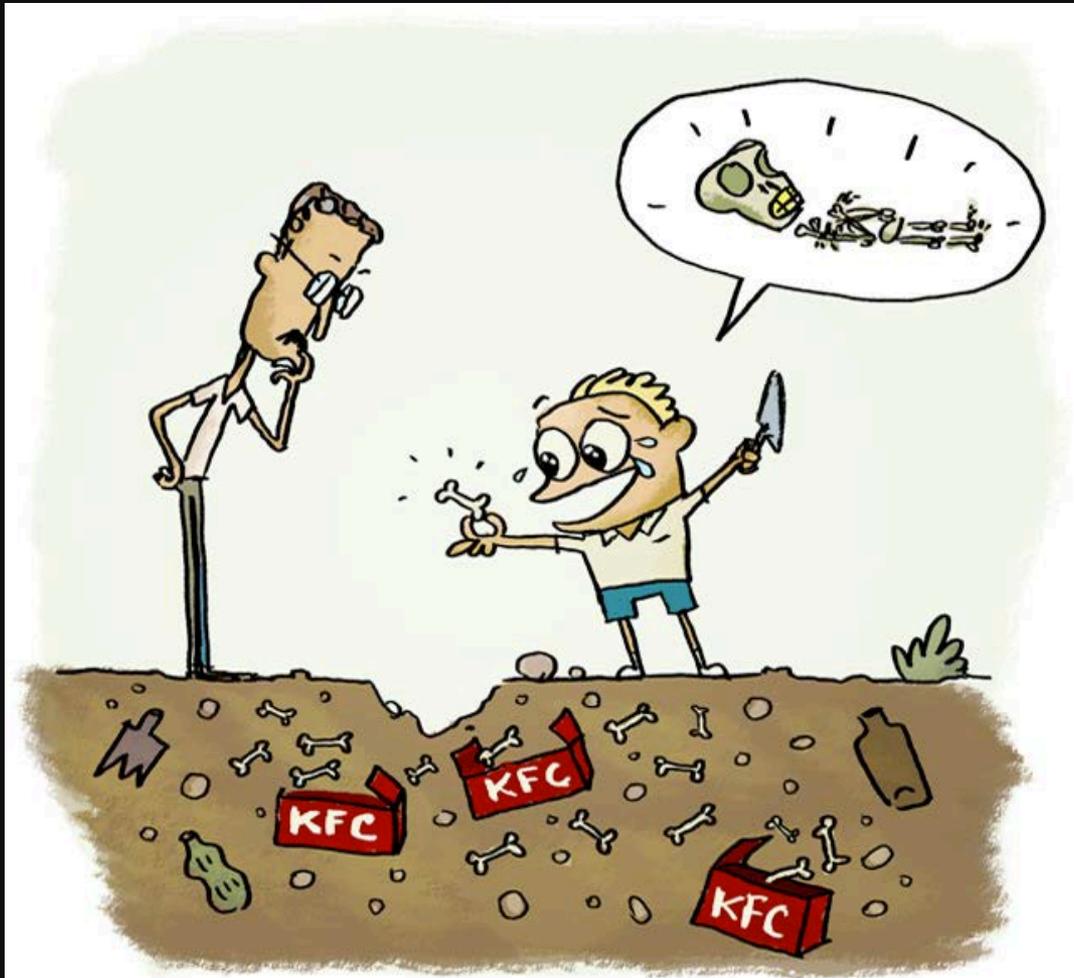
Genomics provides finer scale resolution;
MHI are separate populations from NWHI

Island
O'ahu

Identified pathways of dispersal and origins of
recruitment; management implications



Problems in the field



@PostAntiquarian

Once I got very excited to find some bones in a trench, thinking it may have been a midden, then the KFC wrapper emerged. #fieldworkfail

Problems in the field

- Misidentifications in the field

How many species do you see?



How many species do you see?





Problems in the field

- Misidentifications in the field
- Incorrect labeling



Problems in the field

- Misidentifications in the field
- Incorrect labeling
- Inadequate sample size
- Making sure DNA remains stable
 - Proper preservation solution
 - Avoid extreme heat
- Contamination

Genetics as a tool

- Genetics be can useful in addressing a variety of questions
- Provides insight into evolutionary mechanisms
- Informs management and conservation
- As molecular technology advances, our ability to answer questions increases

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- Castle Foundation
- University of Hawaii Sea Grant Program



NATIONAL MARINE
SANCTUARIES



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Questions?

