



Fish and Aquatic
CONSERVATION

USFWS Fish Technology Centers

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National Fish Hatchery System and Applied Sciences Division

Fish and Aquatic Conservation, USFWS

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Fish and Aquatic Conservation Program



+ 1200
Employees

National Conservation Network



72 National Fish Hatcheries
(1 Historic National Fish Hatchery)



7 Fish Technology Centers



6 Fish Health Centers



1 Aquatic Animal Drug Approval Partnership



51 Fish and Wildlife Conservation Offices

5 Core Programs

- **National Fish Hatchery System**
- Maintenance & Equipment
- Habitat Assessment & Restoration
- Population Assessment & Cooperative Management
- Aquatic Invasive Species

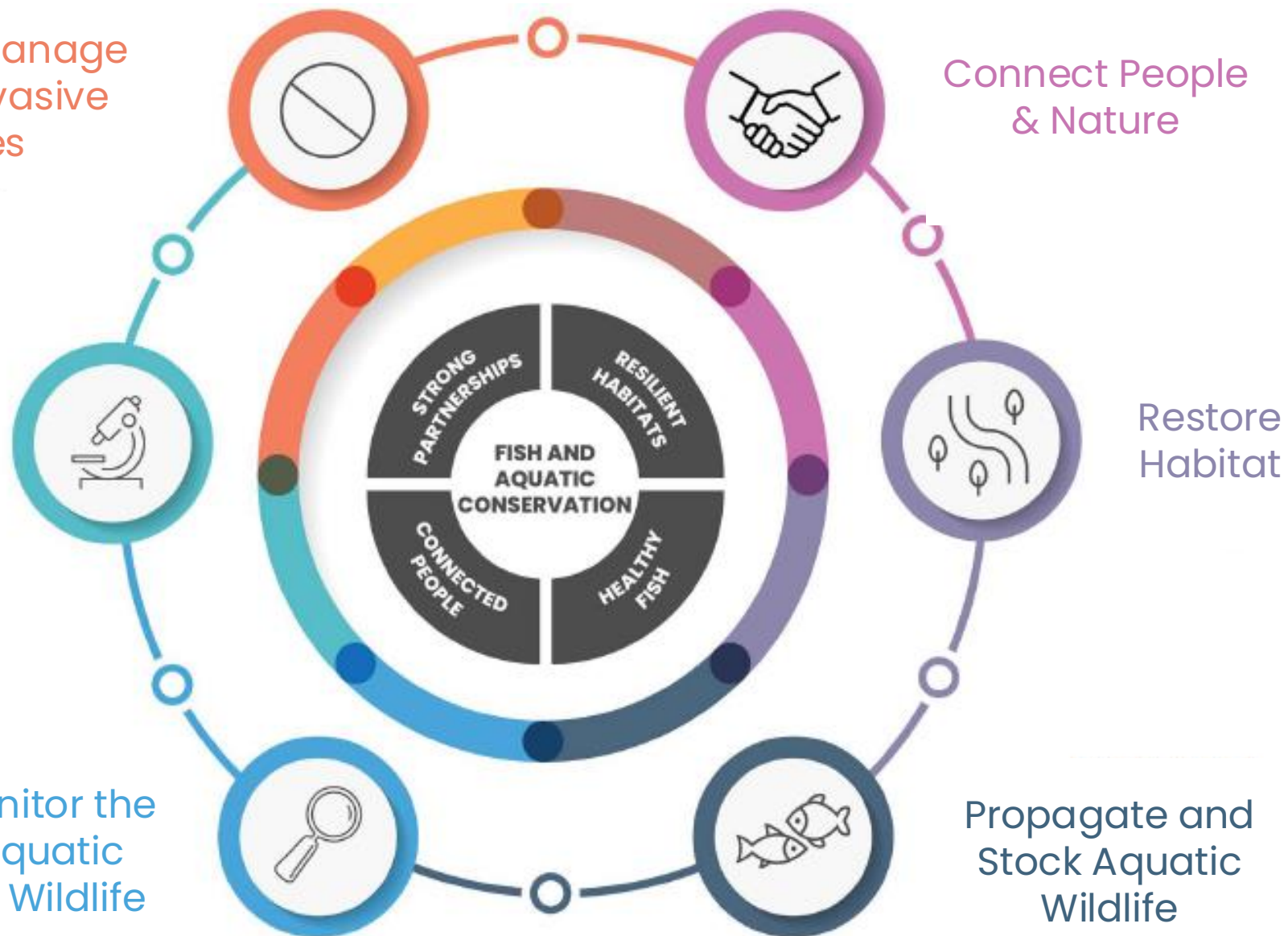


How We Work: Conservation Toolbox

Use & Applied
Science and
Cutting-Edge
Technology

Prevent & Manage
Aquatic Invasive
Species

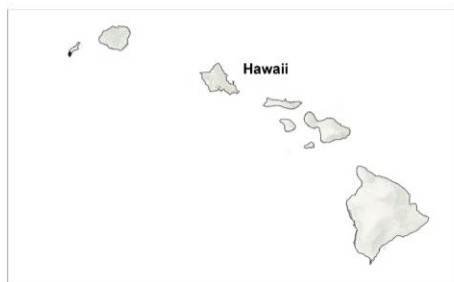
Assess & Monitor the
Health of Aquatic
Habitat and Wildlife






U.S. Fish & Wildlife Service

Fish and Aquatic Conservation Facilities



Facilities

-  Fish and Wildlife Conservation Office
-  National Fish Hatchery
-  Fish Health Center
-  Fish Technology Center
-  Fish and Aquatic Conservation Headquarters & Aquatic Animal Drug Approval Partnership

Research & Applied Science

Mission- Provide leadership in the scientifically based management of national fishery resources through development of new concepts and techniques to solve specific problems in aquatic restoration and recovery activities



Fish Health
Centers



Fish Tech
Centers



Fish and
Wildlife
Conservation
Offices



Aquatic Animal
Drug Approval
Partnership



Fish Technology Center Research Programs

Genetics

Physiology &
Nutrition

Biobanking &
Cryo

Aquaculture
research

Fish Passage

Quantitative
Ecology

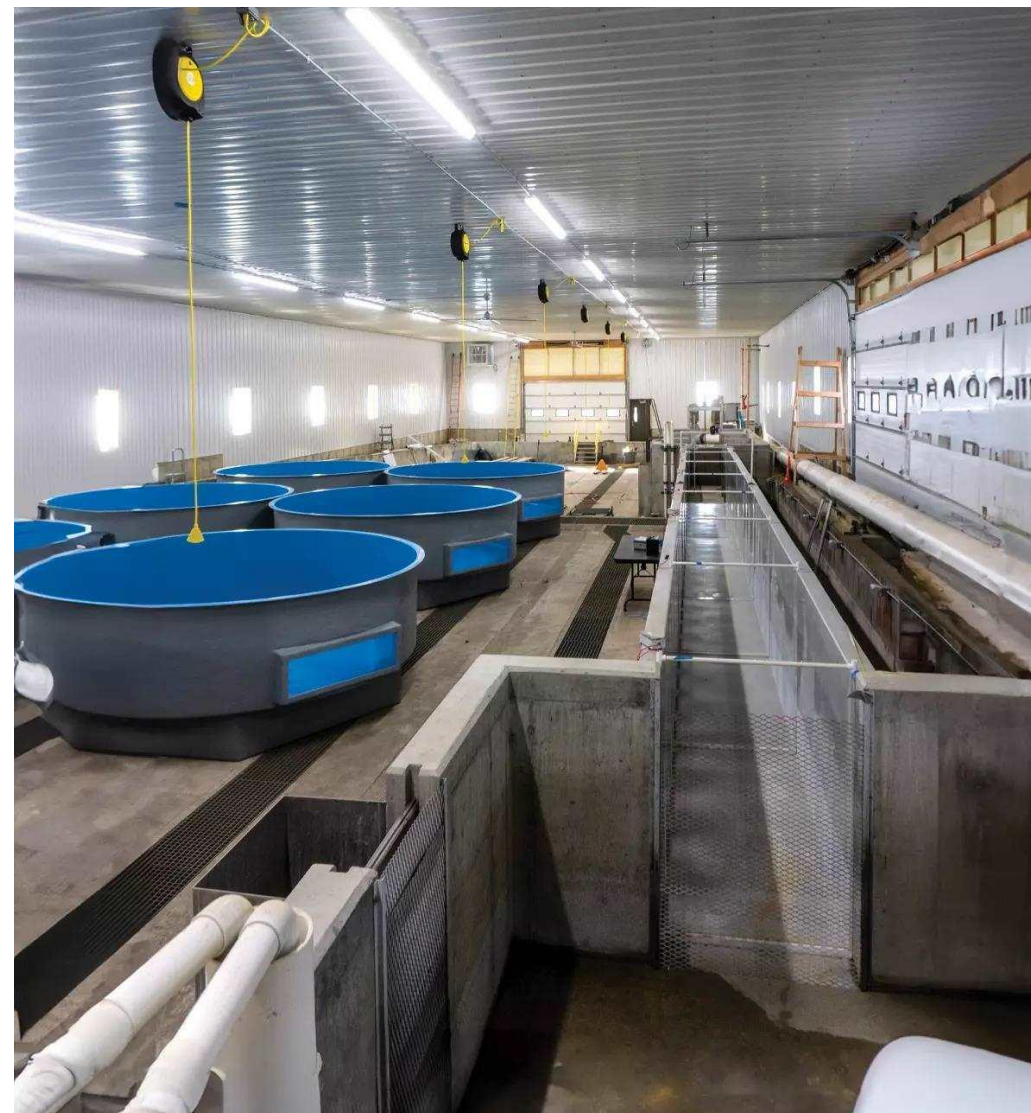
Data
Management

Population
Dynamics

Bozeman FTC

Fish Passage Research Program

- Barrier height
- Flow rate
- Water temp



Abernathy FTC

Quantitative Ecology Research Program

- PIT tag retention in YY male brook trout
- Condition assessments
- New tech development



Assessing superimposition of listed tule fall Chinook salmon redds using aerial and ground surveys on the White Salmon River, Washington

Monitoring & Assessment Work Group

In the Early stages!

- Representatives from all Regions
- Integrates staff across stations
- Opportunity to ID shared needs & priorities
 - Project A & M
 - Biological A & M



Doing
M&A alone



Doing
M&A in
partnership
with NFHP



Research Article

A framework for aquatic invasive species surveillance site selection and prioritization in the US waters of the Laurentian Great Lakes

Andrew J. Tucker^{1*}, W. Lindsay Chadderton¹, Gust Annis², Alisha D. Davidson³, Joel Hoffman⁴, Jon Bossenbroek⁵, Stephen Hensler^{6,7}, Michael Hoff⁸, Erika Jensen⁹, Donna Kashian⁹, Sarah LeSage¹⁰ and Timothy Strakosh¹¹

AIS Invasion Risk

Fish Index (average)

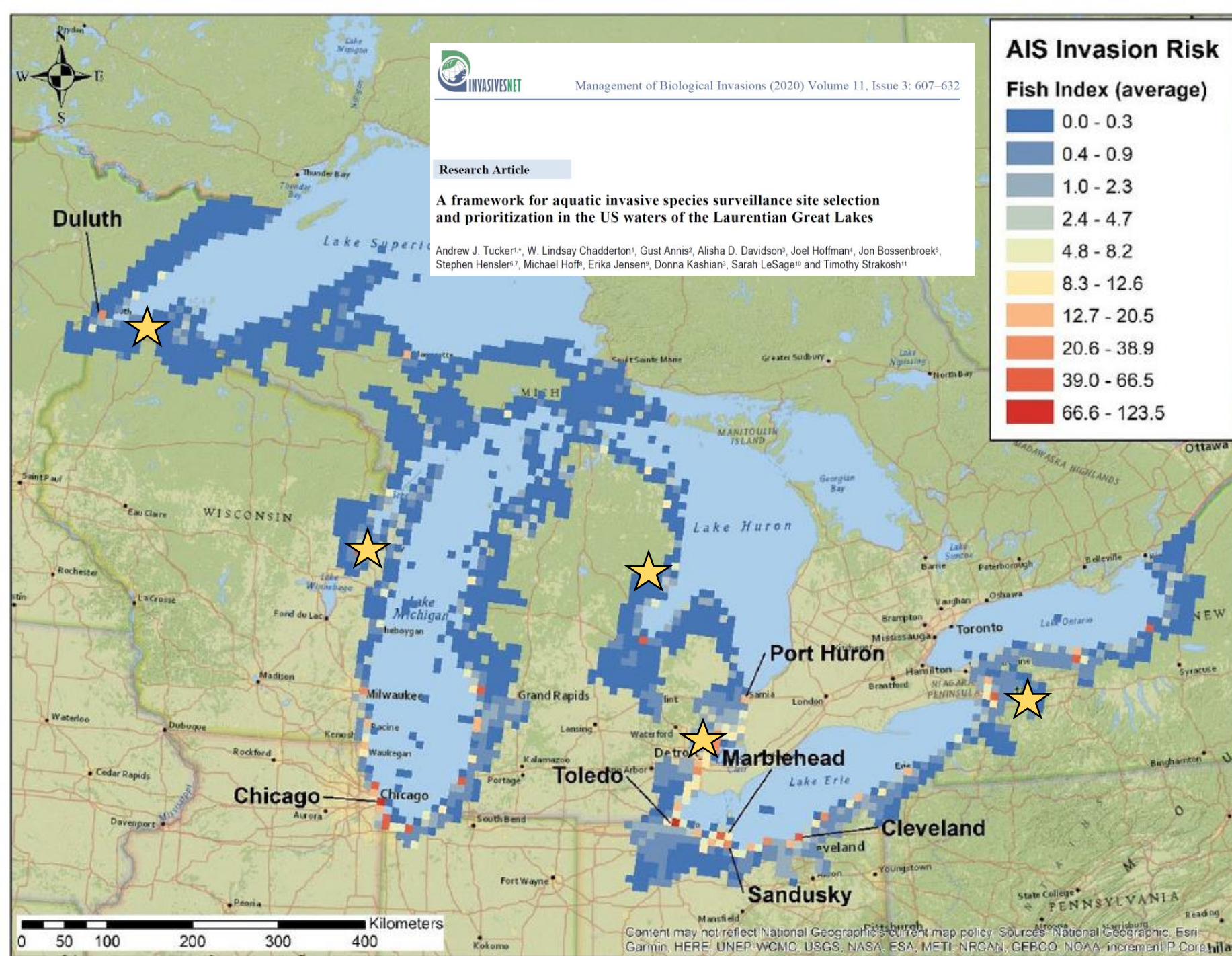
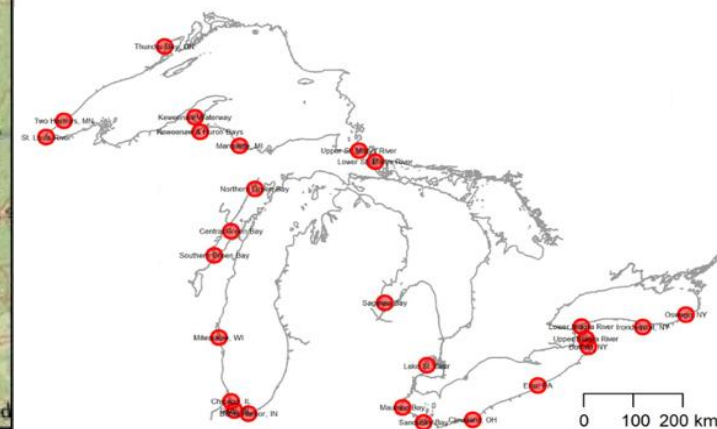


AIS monitoring in the Great Lakes Basin

★ USFWS Fish and Wildlife Conservation Offices

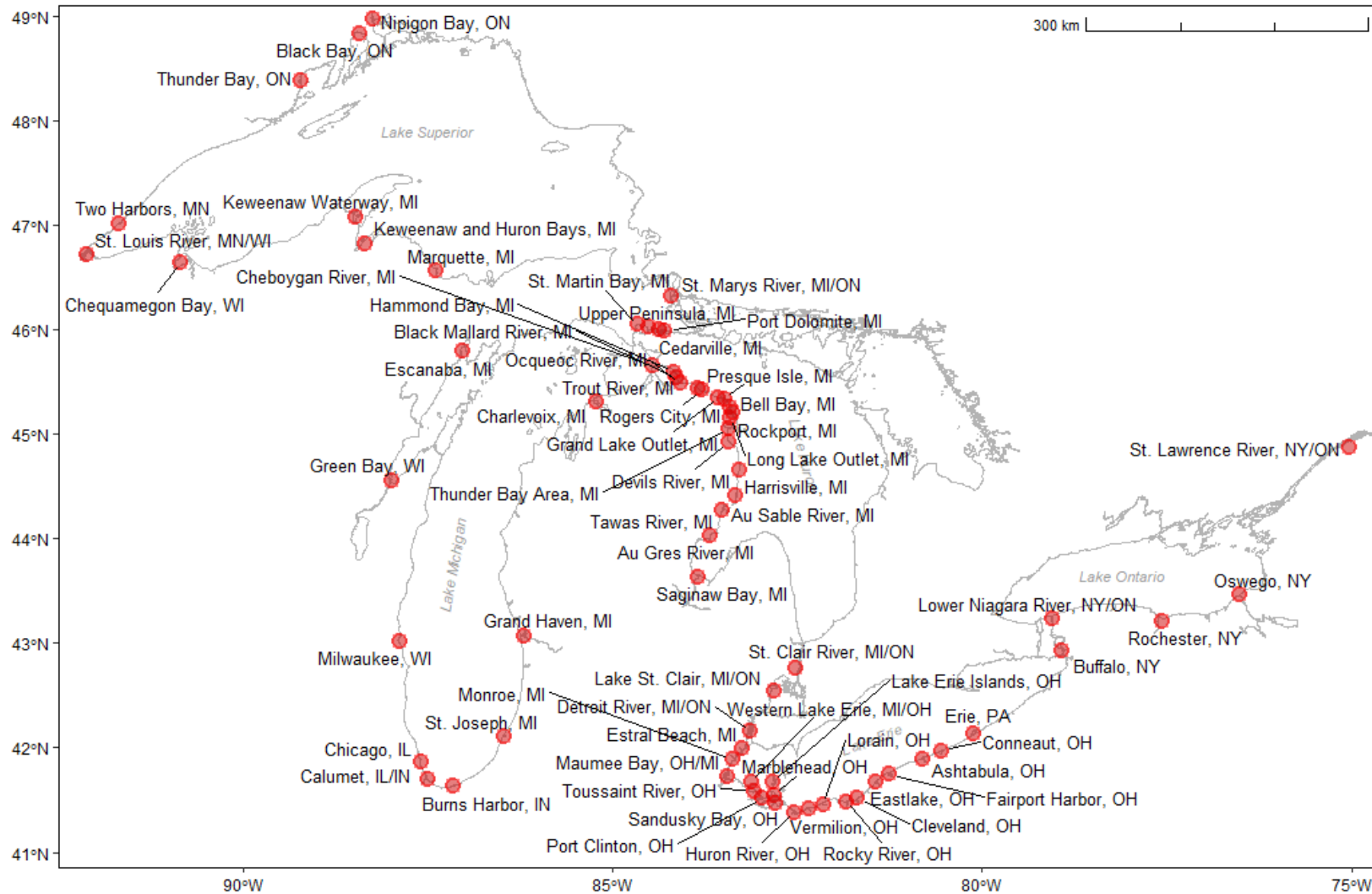
- Ashland, WI
- Green Bay, WI
- Alpena, MI
- Detroit Substation, MI
- Lower Great Lakes, NY

26 hotspots sampled each year for AIS (fish and invertebrates):



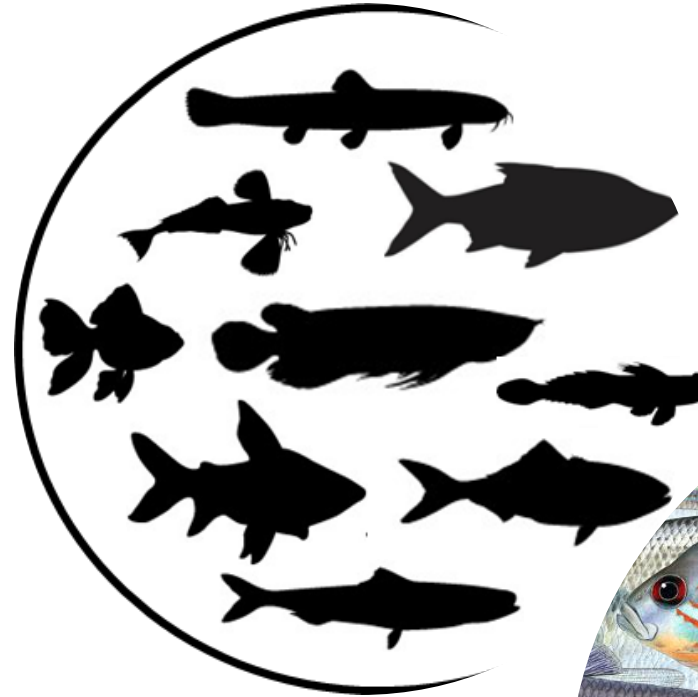
Content may not reflect National Geographic's current map policy. Sources: National Geographic, Esri, Garmin, HERE, UNEP-WCMC, USGS, NASA, ESA, METI, NRCAN, GEBCO, NOAA, incrementIP Corp.

2010–2023 AIS Efforts Monitoring for AIS (fish)



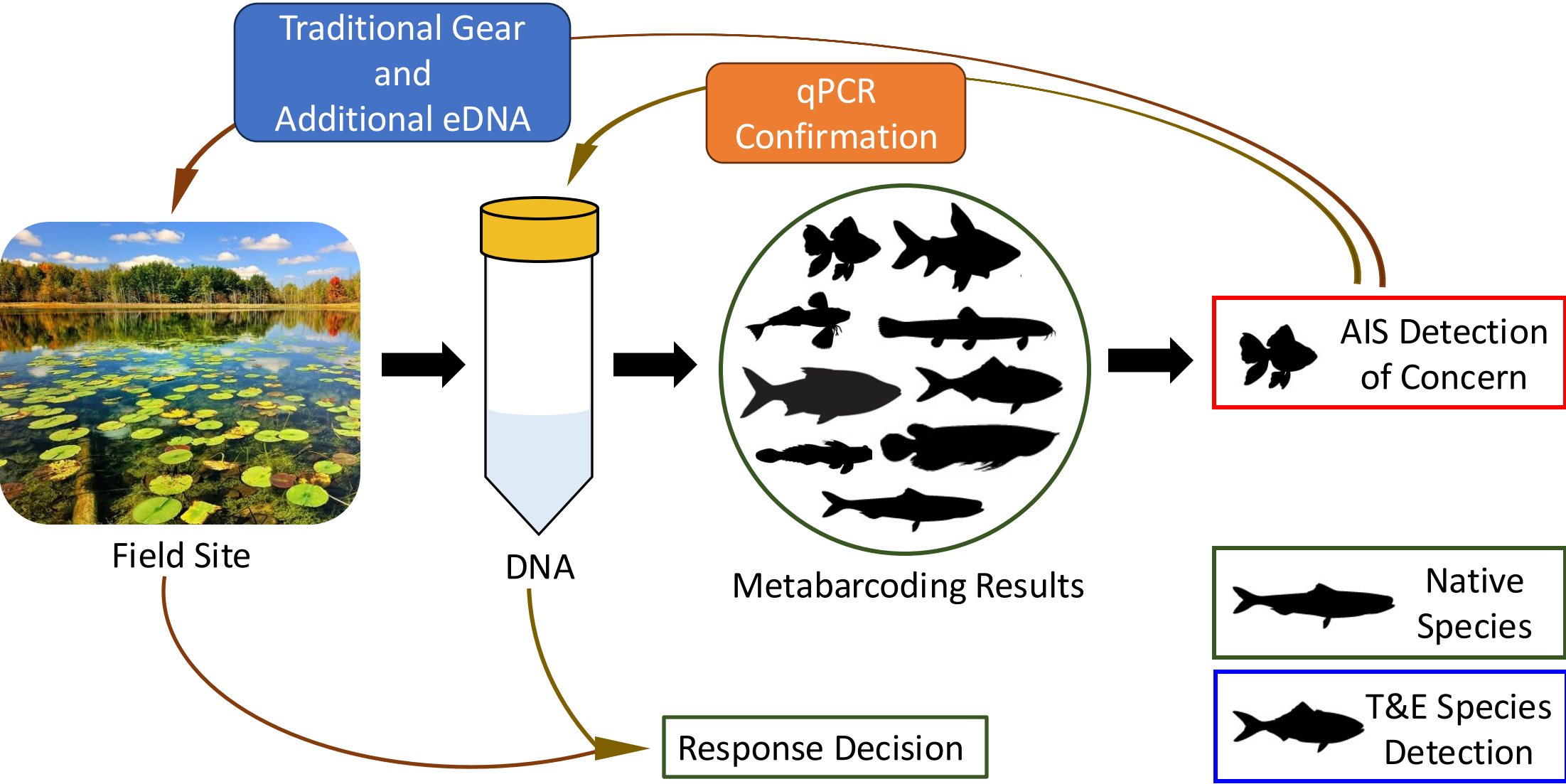
Future AIS Directions

Species of interest is
predetermined –
"Targeted Surveillance"



Species of interest is
undetermined –
"Broad Spectrum Surveillance"

Integrated EDM: Traditional Gear and eDNA Methods

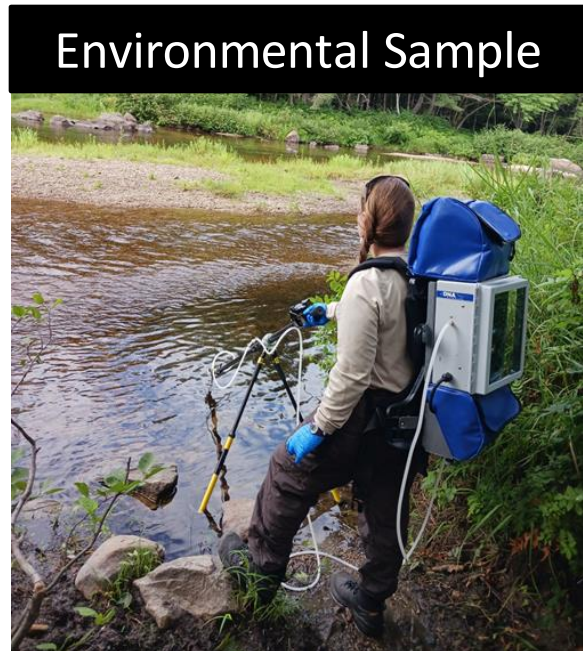


Metabarcoding

Metabarcoding uses generic amplification primers to guide the replication of target DNA and DNA sequencing measure the accumulation of PCR products



Illumina MiSeq DNA Sequencer



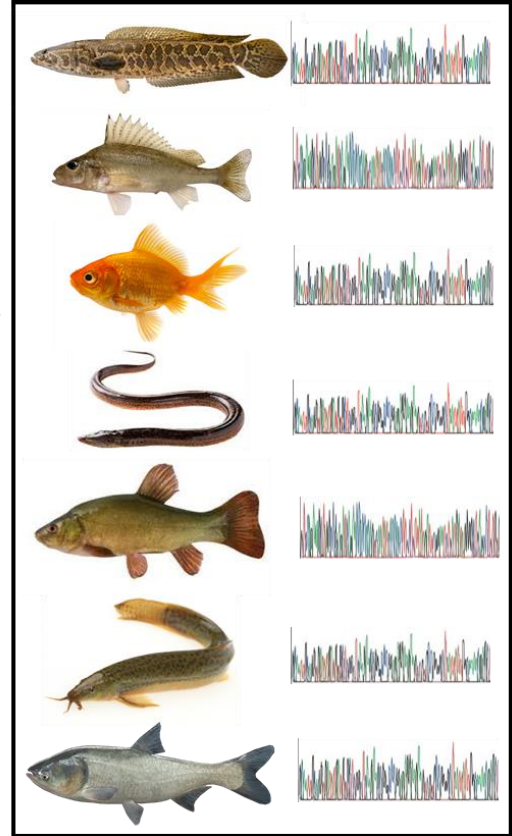
Environmental Sample

DNA sequences recovered from the environment

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AGCTTATTTAGCTGACCTTACTAGAAA
CCTCTGACATAATTTCCCTTCCAGTGGG
ACCTACCCCAATTATTAACCTACACAA
ATTAGATATGCTTGATGATCCCTCGT
GCGGTATTTTAAACCGTGCAAAGGTAGC
CCCTTAATAGCGGCTGCACCATTAGGA
CTACTAGTGCGCCCTACATATGGGA
TACTACGGCTCATACCTTTACAAAGAG
GCACCATTAGGATGTCCTGATCCAACA
ACCCTATGGAGCTTTAGACGCCAACCA
ACATGCCTCAATTGAACCCCGTCCCTI
GTTAATACTCGATTGTTCAATCATCGT
GTGTAGGAACCATATAGGAGTCGAAGC
TCCCTCCCTACGAATCTTTACCTCATI
GAGAAGGGACCTTCCAAGGCCACCACA
GTCCAGGGGTGTGATTCCTGTTGGGGG
ACGTAGGACTTTAATCGTTGAACAAACI
TTAGACGCCAACCAACCACGAAAAGCGI
GATTGTGATTCCAAAGACAGGAGTTCA
GTTCCCTAATGACTCAACGAGGCCAGT
GCCCTACCTAAACAGCCCGTAAAGCCI
GAGTTAGAGGTGGGGTTTCTATGCTTT
ATTTAAAGCAGCCCTTCTTTCTGGGCT
CACCCAGAGTTTTGATCTCTGGAGGAT
GCATAGAAGAAATGAGTCCTAGGGCTAC
GTTAAGCTCTTTATTAAGAACCCTGTC
    
```

Reference database of known species



Read counts – Number of sequences matched

	Sample 1	Sample 2
Bluegill	42931	19793
Alewife	30633	11405
Gizzard shad	499	3385
Pumpkinseed	2467	19162
White sucker	0	994
Bluntnose minnow	0	0
Round goby	127	15551
Northern Snakehead	1395	0
Smallmouth bass	0	1424
Emerald shiner	0	0
Walleye	0	2983
Coregonus	0	98

Reference database is foundational to the metabarcoding effort