Developing Cost Effective Monitoring for Rainbow Smelt using eDNA

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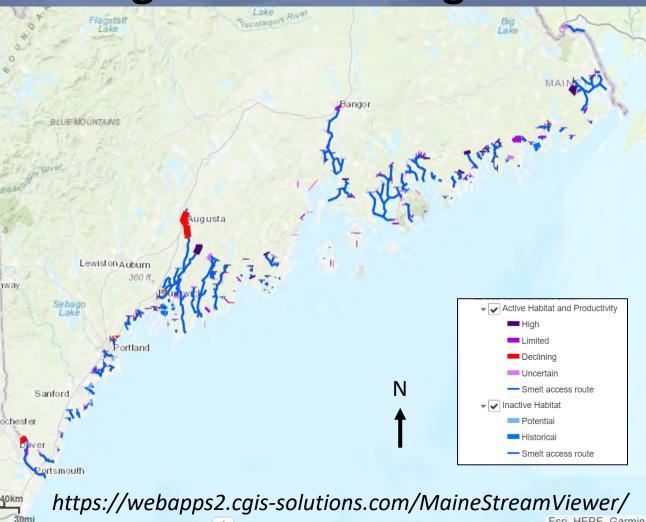
Management Challenges for Rainbow Smelt



- NMFS Species of Concern
- Maine Tier I Species of Greatest Conservation Need
- Declining due to historic alterations to habitat
- ~50% of spawning sites may be impacted by road crossings or dams
- Future climate driven impacts...?
- 2012 Regional Conservation Plan calls for statewide monitoring and restoring access to habitat

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Management Challenges for Rainbow Smelt



- 275 historic spawning stream segments mapped by Maine DMR
- Monitoring so many locations is resource intensive
- Lack of current data hinders restoration efforts

Management Challenges for Rainbow Smelt



Conventional methods are labor intensive or require special training.

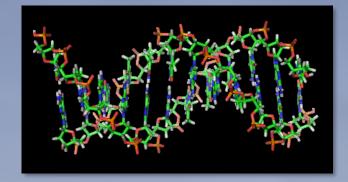
- Trapping
- Trawls

- Creel Surveys
- Egg surveys
- Nighttime observations

eDNA is an Effective and Accessible Tool

Environmental DNA (eDNA): DNA that occurs in an environment as a byproduct of the life processes of living organisms inhabiting that environment or linked environments.

- Less time and labor intensive
- Low risk to smelt populations being monitored
- Does not require special knowledge or equipment
- Highly sensitive, good for rare species
- Presence/Absence information
- Timing and duration of spawning
- Abundance...?
- Biological data collection not possible
- Prioritize locations for conventional monitoring



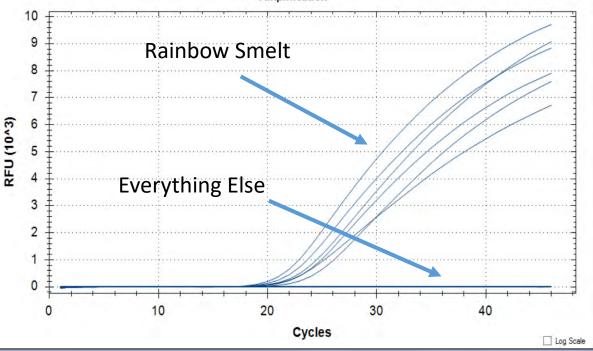
Wikipedia.org

eDNA is an Effective and Accessible Tool

Smelt TaqMan MGB-NFQ qPCR Primer-Probe Set develop by Kinnison Lab, University of Maine

OSM ND5 Primer-Probe Set

Species	5' Forward 3'	5' Probe 3'	5' Reverse 3'	
OSM	CACAACCATCCACCGCTCTCTC	TGGGCCAAGCTATCGCGAGCCA	TTGCTGCTGGGAGGTGAATAGC	
ARC	TGCCGTCGTCCACCGATTAGCC	TAGGACAAACTATTGCCAGCCA	T <mark>GA</mark> CT <mark>ATG</mark> GG <mark>C</mark> AGGT <mark>T</mark> A <mark>G</mark> T <mark>T</mark> GA	
ATL	AGCCATTATCCACCGATTAGCC	TAGGACAAACCATTGCCAGCCA	T <mark>GA</mark> CCATAGGTAGGTGAGTGA	
BKT	TGCCATCGTACACCGATTGGCC	TAGGACAAACTATTGCCACCCA	T <mark>AA</mark> CT <mark>ATA</mark> GG <mark>G</mark> AGGT <mark>T</mark> A <mark>G</mark> TTGA	
LKT	TGCCGTCGTCCACCGATTAGCC	TAGGACAAACTATTGCCACCCA	T <mark>GA</mark> CT <mark>ATG</mark> GG <mark>C</mark> AGGT <mark>T</mark> A <mark>G</mark> TTGA	
RBT	C <mark>GCC</mark> A <mark>T</mark> CATCCACCG <mark>A</mark> T <mark>TGAC</mark> C	T <mark>A</mark> GG <mark>A</mark> CAA <mark>A</mark> CCATCGCCAGCCA	T <mark>GA</mark> CT <mark>AT</mark> TAG <mark>GC</mark> AGG <mark>TGAG</mark> TTG	
BNT	AA <mark>CC</mark> A <mark>T</mark> CATCCACCG <mark>A</mark> TTGGCC	TGGG <mark>A</mark> CAA <mark>A</mark> CCAT <mark>T</mark> GC <mark>T</mark> AGCCA	T <mark>GA</mark> CCATA <mark>GGC</mark> AGGTGA <mark>GTT</mark> GA	
LWF	C <mark>G</mark> C <mark>ATTG</mark> TCCACCG <mark>ACTAGC</mark> C	TCGGTCAGGCCATTGCCAGCCA	T <mark>AAT</mark> T <mark>ATG</mark> GG <mark>C</mark> AG <mark>A</mark> TG <mark>GG</mark> T <mark>G</mark> GA	
СР	ATCAATTATCCATCGGATAATT	TGGG <mark>T</mark> CAA <mark>AAA</mark> AT <mark>T</mark> GC <mark>TAC</mark> CCA	T <mark>AATAA</mark> GGGGTAGATTAATTGA	
NP	ATCAATTATTCATCGCTTAACC	T <mark>C</mark> GGCCAA <mark>AAA</mark> AT <mark>T</mark> GC <mark>C</mark> ACCCA	T <mark>AATTATG</mark> GGGAGGT <mark>TG</mark> AT <mark>G</mark> GA	
LMB	AGGGGTTTTTCCACCGCCTCATG	T T GGCCAAGC <mark>A</mark> AT T GC <mark>C</mark> AGCCA	AAGCTAAGGGGGGTATTGAGGGA	
SMB	CA <mark>T</mark> A <mark>GTTT</mark> TCCACCGC <mark>CT</mark> T <mark>A</mark> TG	TTGGCCAAGC <mark>A</mark> ATTGC <mark>C</mark> AG <mark>T</mark> CA	AAACTAAAGGGGTGTTAAGGGA	
BC	AATAATCACTCACCGCTTCATT	T <mark>A</mark> GGCCA <mark>GT</mark> C <mark>A</mark> AT <mark>T</mark> GCTAGCCA	T <mark>AAT</mark> TAAG <mark>GGGGT</mark> GT <mark>TG</mark> A <mark>GG</mark> GA	



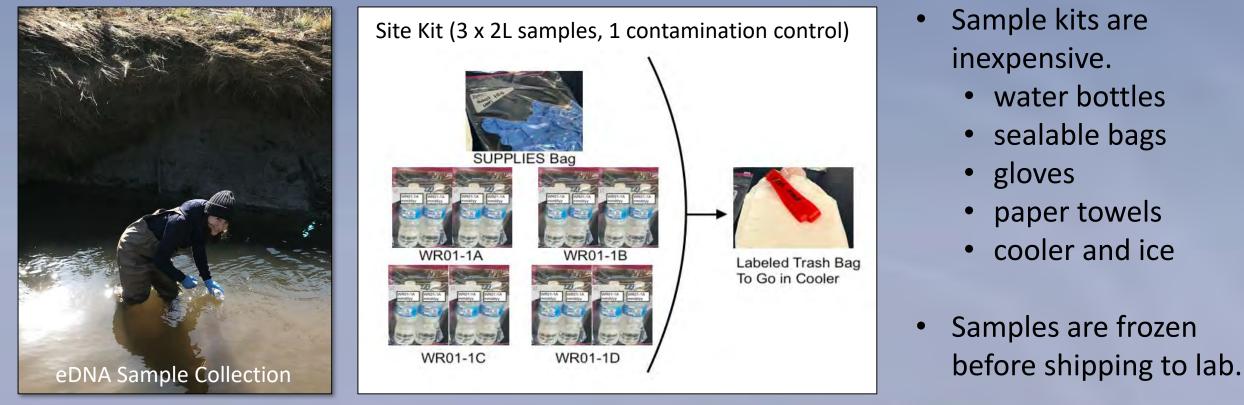
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Amplification

+ Initially Field Validated by Testing Positive in Archived Floods Pond Water

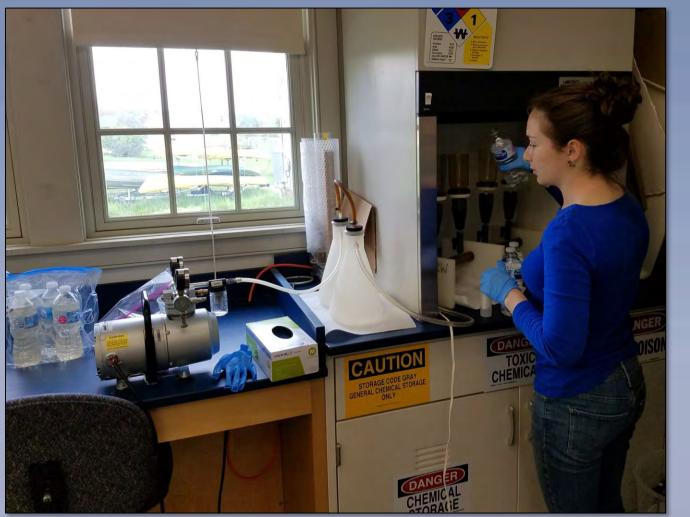
eDNA is an Effective and Accessible Tool

eDNA surveys could in principle be conducted by almost anyone, without risk of harm to protected species, without concern for legal harvest seasons, and without special licenses or permits.



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eDNA is an Effective and Accessible Tool



Detailed protocols have been developed.

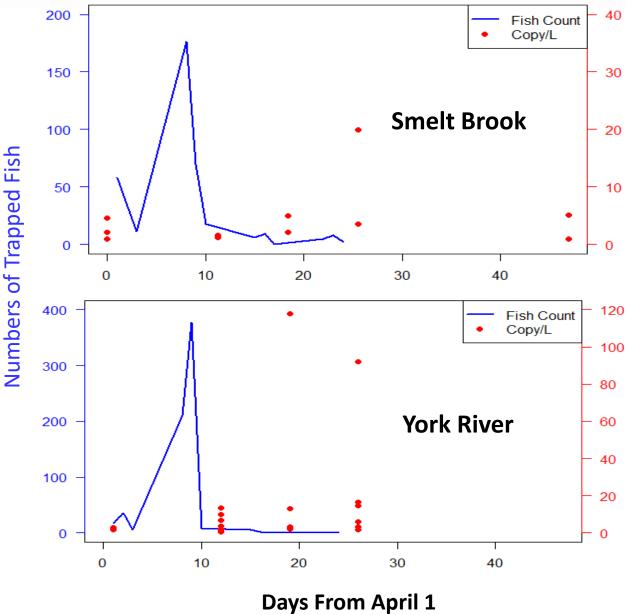
- EPA approved mini-QAPP
- UMaine sampling protocols and filtering protocols

Filtration in-house saves shipping and lab costs, but contamination control is a major consideration.

Monitoring in partnership with qualified lab.

Smelt eDNA Pilot Study

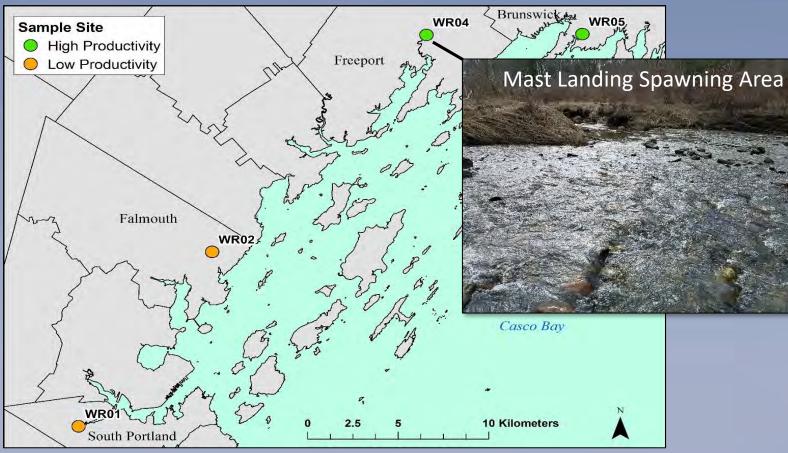




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eDNA Copies Per Liter

2018 Casco Bay Study



- Sampling March 29 to May 9
- Downstream from spawning areas
- 2 liter samples collected at three locations (A, B, C)
- 177 samples total

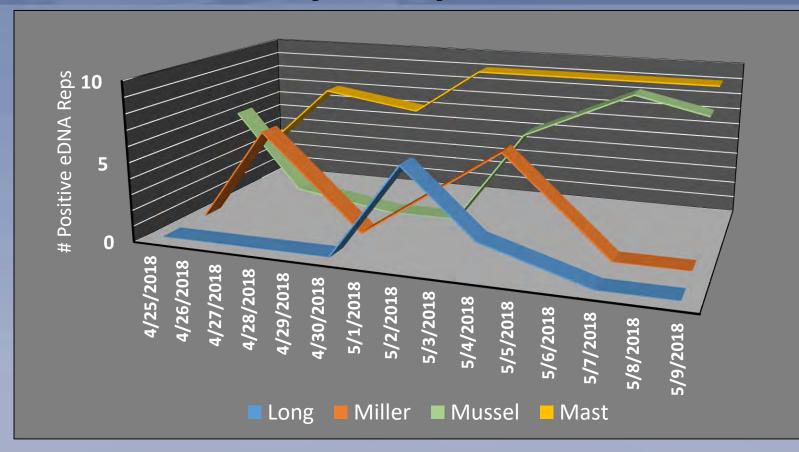
Casco Bay study sites at:

- Long Creek (WR01)
- Mussel Cove Creek (WR02)

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- Mast Landing (WR04)
- Miller Creek (WR05)

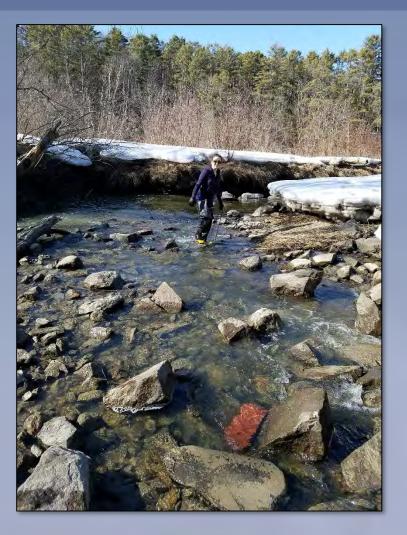
2018 Casco Bay Study



- Unlike pilot study, initial lab detection was poor
- Environmental inhibitors reduced amplification
- PCR inhibition clean-up increased reliability
- eDNA is able to detect rainbow smelt through time and across a range of abundances

Next Steps

- Calibrate for environmental conditions (flow, inhibition, breakdown rates)
- Further study of eDNA dynamics (sampling for eggs vs. adults)
- Develop application of techniques for managers and restoration planners
- Explore possibility of engaging citizen scientists in surveys



References

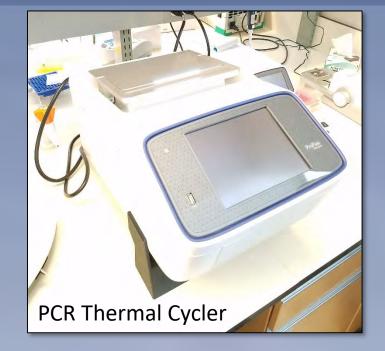
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Thank You!

Project Partners

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US Environmental Protection Agency Maine Outdoor Heritage Fund