



Northern Cod Fisheries Improvement Project:

Improving the Future through Improved Science

BARCELONA APRIL 2023

ASP/AGC Northern Cod FIP – Our Objectives

 Address gaps in scientific information to facilitate sustainable management of this important resource

Prepare information and documentation for eventual MSC certification

Partners



- DFO and Govt NL
- Important International Customers
 - Marks and Spencer (M&S)
 - Sysco France
 - High Liner Foods
 - · Young's Seafood
- Sustainable Fisheries Partnership
- Academia
 - Ocean Tracking Network
 - Dalhousie University
 - Memorial University
- Inshore Processors + Year-round Harvesters
 - ASP Members
 - AGC Members

FIP at a Glance

FIP at a Glance		
ENVIRONMENTAL ① Addressing 28 of 28 indicators Current Status:		
7%	93%	
Starting Evaluation: February 01, 2015 11% 32% 57%		
Progress Rating 1	Actions Complete 1	Next Update Due () JUN 2023
A	91%	Target End Date
ADVANCED PROGRESS	Complete Incomplete	DEC 2025
Additional Impacts:		
SOCIAL ()		
Risk Assessment 1 : Not Required		

- FIP started in 2015
- 13 FIP tasks completed
- 3 Outstanding Action Items
 - Stock Status monitoring Annual
 - HCR testing To be completed during Rebuilding Plan review.
 - Northern Cod Acoustic Tracking (NCAT) Project – Stock migration and composition research

ASP/AGC FIP Working Group: Data Collection and Research Continues

Northern Cod Acoustic Tracking (NCAT) Project Update

- NCAT Project Elements
 - Research plan to investigate Northern Cod migration & stock composition
 - Genetic studies provided insight into genes linked to behavior, habitat choice and diet
 - Objective to acoustically tag 1200 cod & track migrations on Newfoundland Shelf.
 - 75 receiver acoustic array installed in 2020 →
 - 775 cod tagged to date
 - Receiver data uploaded via autonomous Wave Gliders



2019 Tagging Results



- July/ August 2019
- 338 acoustic tagged & genetically sampled cod

 Tagging conducted in 2J, 3K and 3L



2021 Tagging Results

- May 2021
- 66 cod tagged and genetically sampled, (51 in 2J, 15 in 3K)
- 49 additional cod externally floy tagged





March 2022 Tagging Results



- 371 acoustic tagged & genetically sampled cod
- 298 Floy tagged cod, photographed for morphological body shape analysis
- 25 fishing sets
- 14 recons to find cod aggregations
- Tagging team from HusseyLab at the University of Windsor and Fisheries and Oceans Arctic region / the University of Manitoba



2023 Spring Tagging Trip

- April 2023
- 7-day trip into 3L and 3K
- Bottom temps were colder
- Observed capelin, jellyfish, flatfish
- Unfortunately, no suitable cod found or tagged.



Initial Results!

Initial Genomic and Telemetry Results!

- MITACS funding to complement salary of postdoctoral fellow.
- Continue data analyses to integrate telemetry results with genomics.
- PhD Lisette Delgado
 - Postdoctoral fellow at Dalhousie University, currently working on Northern Cod.
 - Biologist, specialized in population genomics and bioinformatics.
 - M.Sc. studies, Michigan State University & Ph.D. studies, Dalhousie University.
 - Proficient in molecular lab techniques and data analyses in the fields of genomics (reduce representation and whole genome) and transcriptomics.
 - Skilled in programming languages: bash, R, and python.
 - Published in top tier journals (ecology and evolution fields).

Previous NCAT Genetics Studies vs Whole Genome

2019 & 2020 genetics studies on Northern Cod

- · Provided valuable information on genetic make up
- Analyzed 12K 25K markers using reduced representation SNPs
- Coarse assessment indicated genetically similar population
- Example:

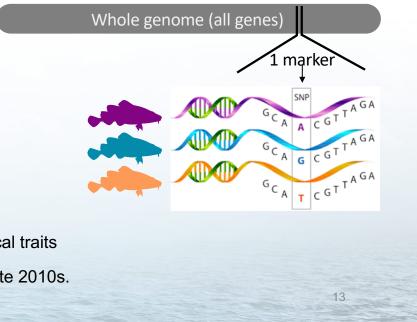
REGULAR PAPER

JOURNAL OF **FISH** BIOLOGY

Life-stage-dependent supergene haplotype frequencies and metapopulation neutral genetic patterns of Atlantic cod, *Gadus morhua*, from Canada's Northern cod stock region and adjacent areas

Gregory Neils Puncher^{1,2,3} | Sherrylynn Rowe³ | George A. Rose⁴ | Geneviève J. Parent² | Yanjun Wang⁵ | Scott A. Pavey¹

- Current work:
 - Whole genome approach (complete set of cod genes)
 - Analyses > 1 million markers
 - Fine-scale differentiation looking at all genes for biological traits linked to migratory, spawning and other characteristics
 - Temporal analysis: samples from the early 1990s and late 2010s.



Preliminary Results Summary

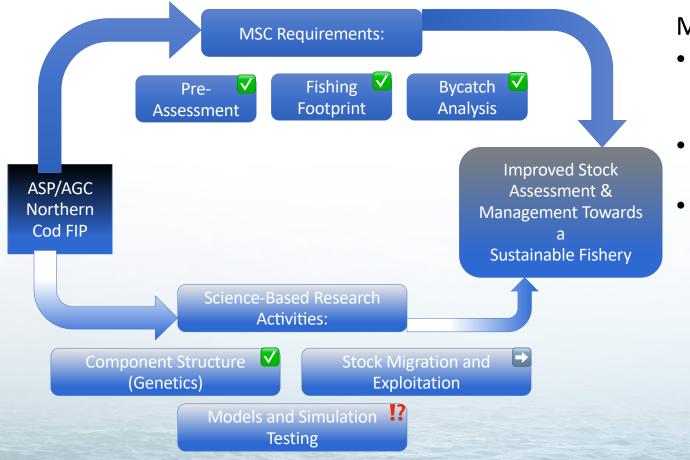
Genomics

- Identified >1.5 million markers
- Inshore Labrador individuals are the most genetically distinct
- Genetic differentiation in 2J3KL is driven by two supergenes, these variations might be associated with migratory behavior and spawning time.

Telemetry (tracking acoustically tagged cod)

- >300,000 detections
- 378 individuals were detected at least once after their release
- Individual movement can be classified:
 - Wide distribution (move between 2J, 3K, 3L, and 4R divisions)
 - Narrow distribution (remain within one division)
- ~20% of NCAT tagged individuals seem to remain only at inshore locations
- Remaining ~80%, spend the winter and spring at mostly at offshore locations.
- Too soon to estimate what percentage remain offshore throughout the year.

ASP-AGC FIP Roadmap – What's next for NCAT?



May 2023

- Convene scientists and stakeholders to discuss:
- Summer 2023 tagging approach
- Next phase of NCAT project:
 - Data analysis
 - Expected projects
 - Incorporation in stock assessment process

15

Questions?

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