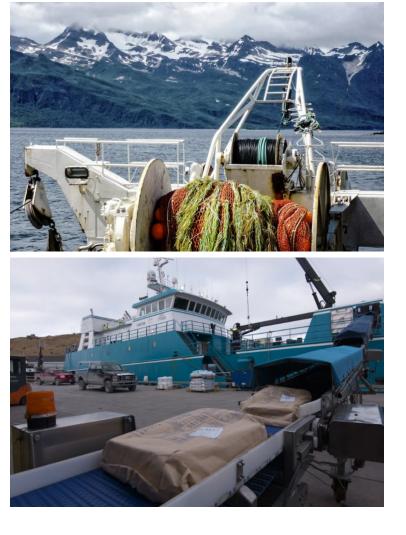
Molecular indicators of performance in a marine fish

exposed to temperature stress



Laura Spencer, NOAA AFSC Affiliate University of Washington SAFS & CICOES



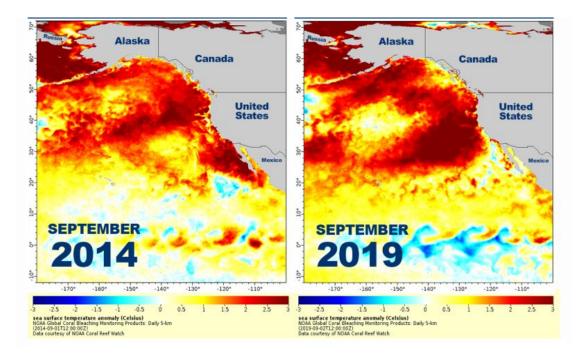
NOAA surveys & manages Alaskan fisheries (AFSC)

VALUE & VOLUME OF KEY SPECIES, 2019

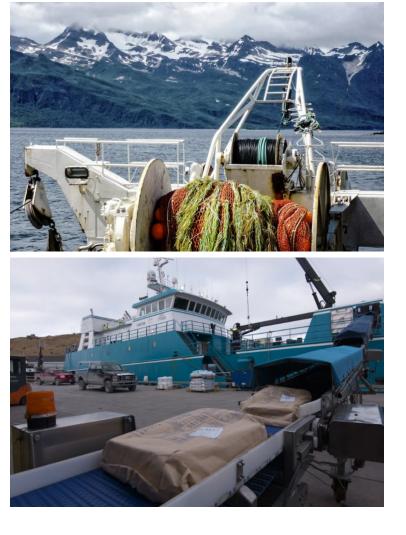


ASMI (industry report), 2022

Heatwaves!



NOAA feature story, September 05, 2019



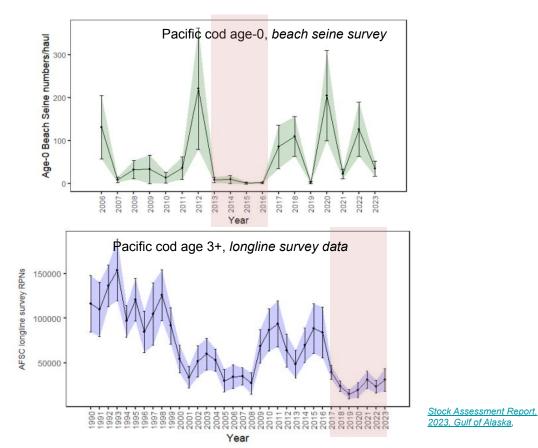
NOAA surveys & manages Alaskan fisheries (AFSC)

VALUE & VOLUME OF KEY SPECIES, 2019





Pacific cod, Gadus macrocephalus



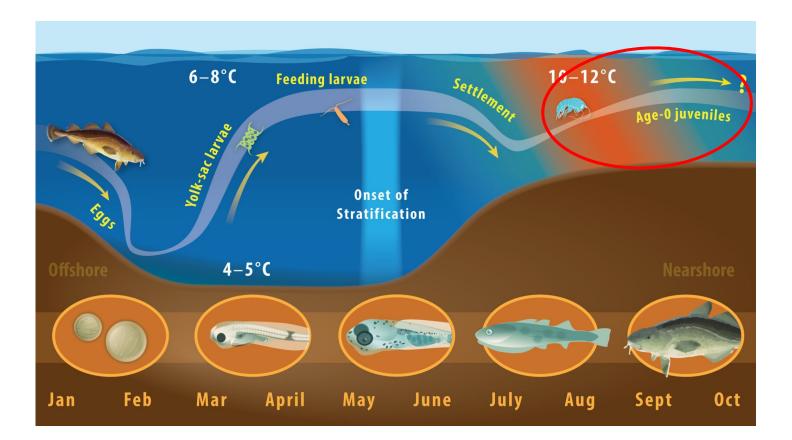


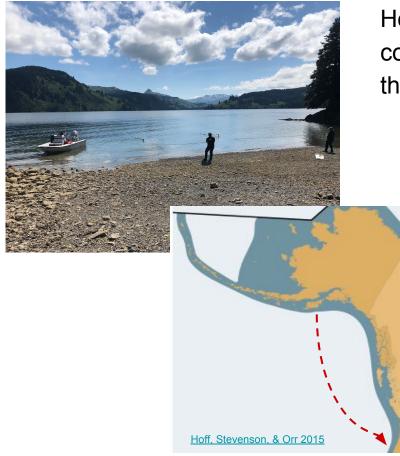


Big Questions

- Why & how does warming affect Pacific cod recruitment?

- Do stock assessment reference points need to be updated?
 - E.g. natural mortality rates
- How resilient are Pacific cod populations in Alaska to warming?





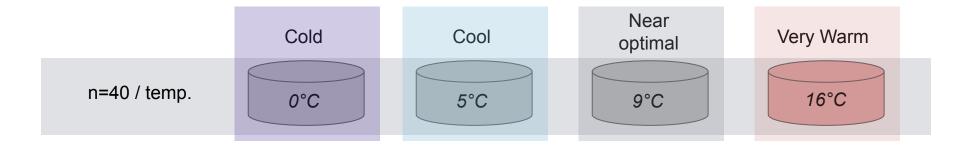
How does temperature affect **juvenile** Pacific cod **growth**, **survival**, **& energy allocation** in the pre-winter period?

- Wild juveniles (age-0) caught off Kodiak, AK late summer
- Transported to Newport, OR wet lab
- ~6 weeks experiment (age-0)



How does temperature affect **juvenile** Pacific cod **growth**, **survival**, **& energy allocation** in the pre-winter period?

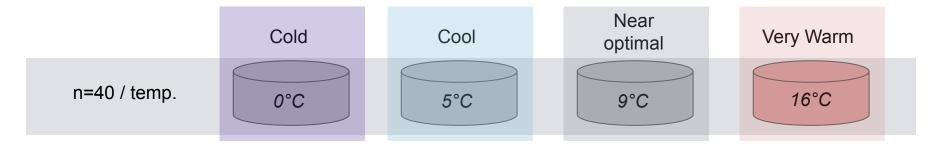
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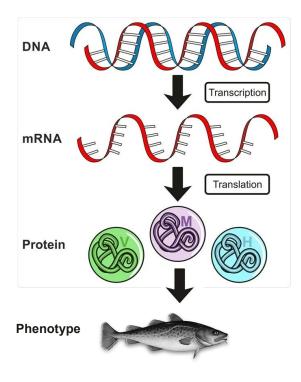


Individuals tagged, collected:

- a. Genetics with fin clips, n=40/temp (lcWGS)
- b. Growth rates (length & wet weight) during acclimation, treatment
- c. Body condition (Kwet)
- d. Liver condition (HSI)
- e. Survival
- f. Liver lipid components (n=25/temp)
- g. Gene expression with liver, n=18/temp (RNASeq)



"Genome-to-Phenome" dataset, Pacific cod juvenile temperature response



Adapted from: udaix/Shutterstock.com

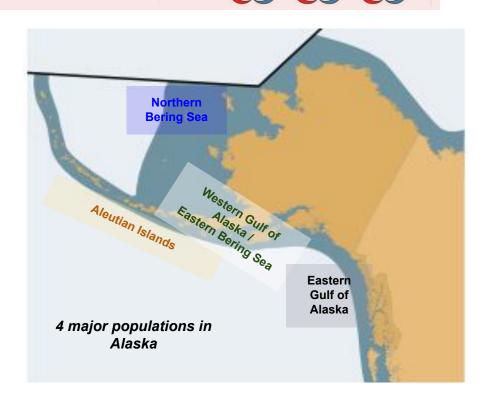
- 1. Genetics Who are they?
- 2. Phenotypes How does warming affect key traits?
- Integrate datasets –
 Why are some fish less sensitive?

"Genome-to-Phenome" dataset, Pacific cod juvenile temperature response

1. Genetics – <u>Who</u> are they (wild caught)?

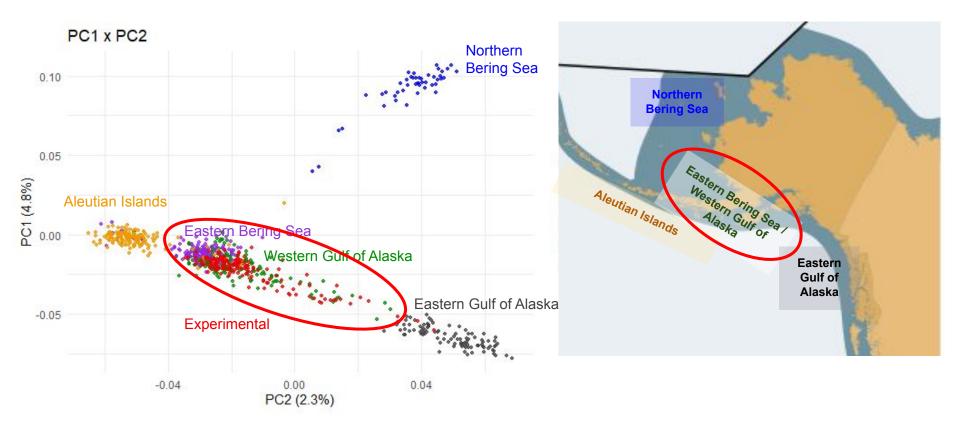
Tools used:

- Sequence whole genome ~3x (i.e. "low-coverage"), n=160
- ~\$100/fish, genome size ~500Mb (similar to pacific oyster)
- High-quality reference genome for alignment & genotype data (bwa mem & angsd)
- <u>More</u> data from ~55 reference fish per population, assign our fish (wgsassign)
- 6,101 sites used to predict population of origin of experimental fish, 96% accurate



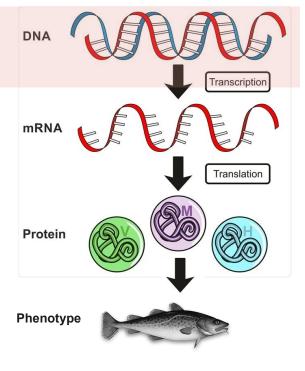
DNA

Genetics - They are the Western GOA / Eastern Bering Sea group



"Genome-to-Phenome" dataset, Pacific cod juvenile temperature response

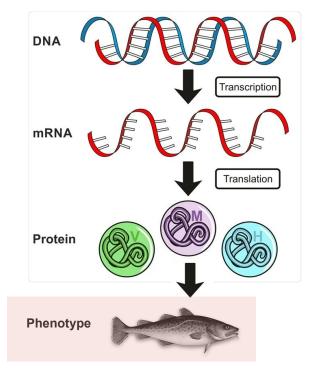
Genetics – they are one population
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Adapted from: udaix/Shutterstock.com

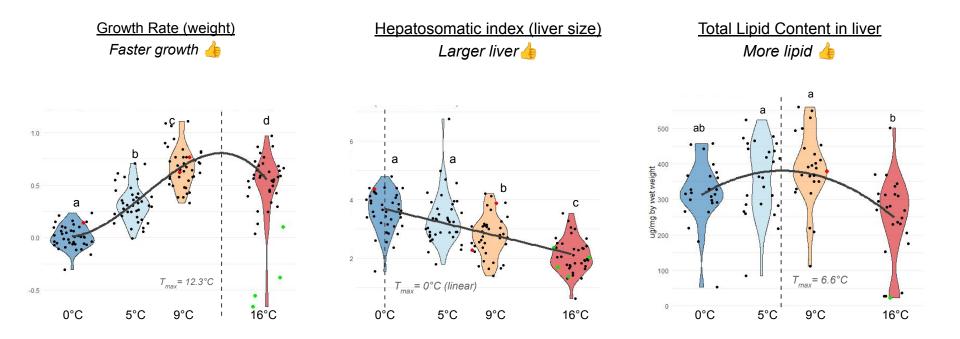
"Genome-to-Phenome" dataset, Pacific cod juvenile temperature response

- Genetics they are one population
 western Gulf of Alaska / Eastern Bering Sea
 - Phenotypes <u>how</u> are survival-associated biometrics affected?



Adapted from: udaix/Shutterstock.com

Warming decreased lipid reserves

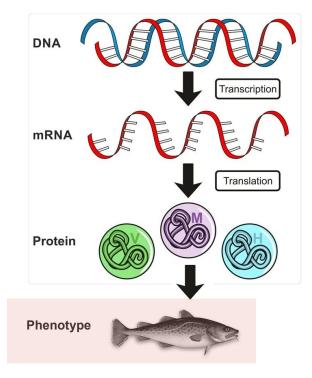


Survived

Died

"Genome-to-Phenome" dataset for juvenile Pacific cod

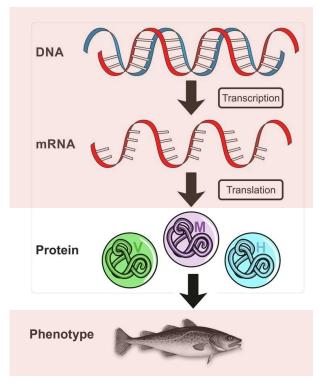
- Genetics they are one population
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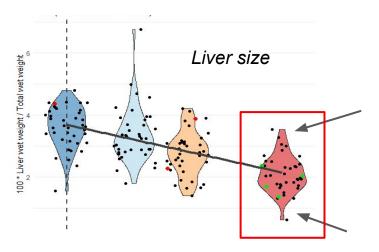
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 - Integrate datasets Performance indicators!
 - a. Genetic variants
 - b. Expression patterns



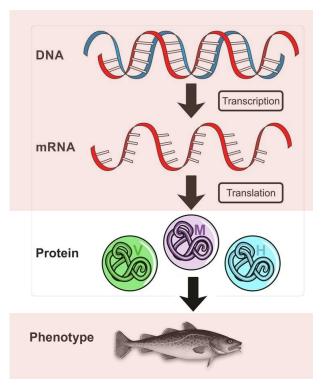
Adapted from: udaix/Shutterstock.com

Variation within each temperature - opportunity!



- 1. Identified <u>sites on genome</u> associated with liver size, lipid content, & growth in warming
- 2. Identified best sites/genotypes that predict performance in warming, are adaptive

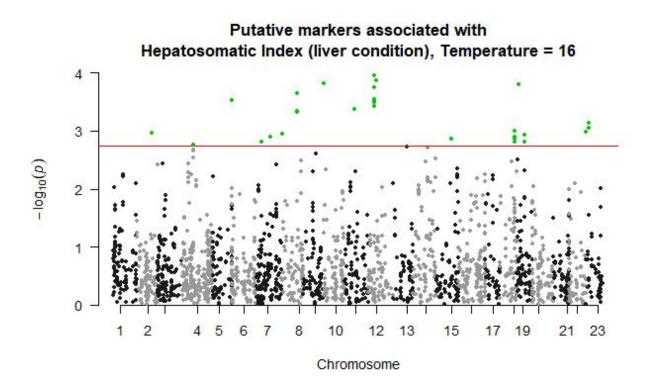
Genome-Wide Association Studies (GWAS) within each treatment



Adapted from: udaix/Shutterstock.com

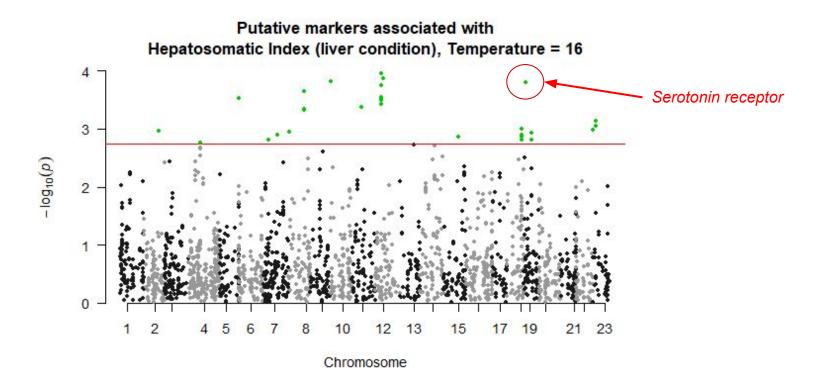


~100 markers putatively associated with liver size in Pacific cod juveniles exposed to warming



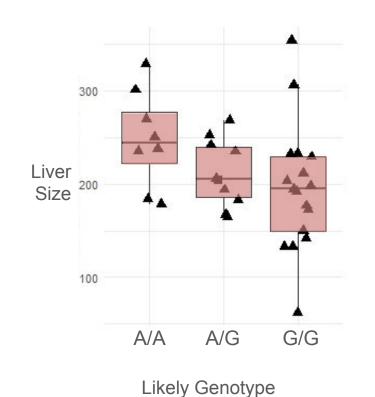


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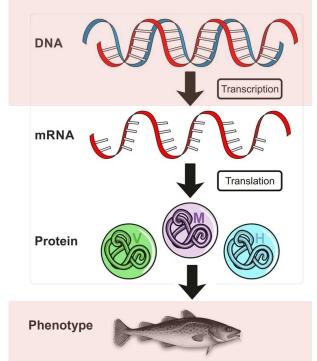


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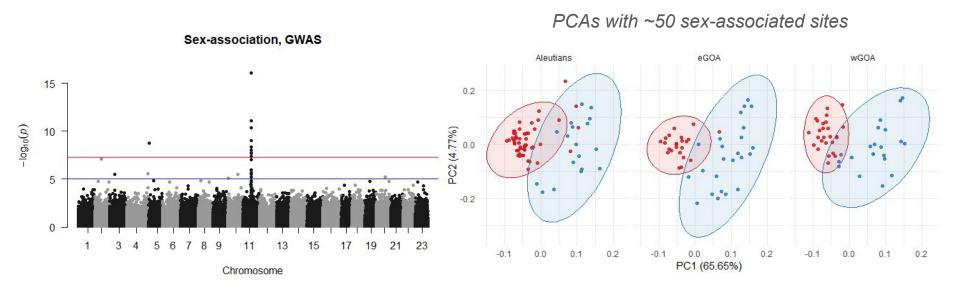
Marker in a gene coding for a **receptor for serotonin** (5-HT4) which regulates appetite "Genome-to-Phenome" dataset for juvenile Pacific cod

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- 1. Integrate datasets Performance indicators!
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Adapted from: udaix/Shutterstock.com

Example Genome Wide Association Study – Sex markers discovery (preliminary)



Female Male

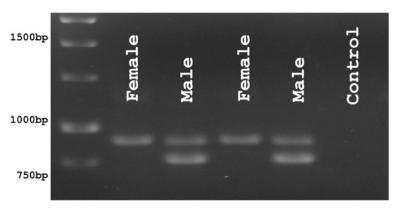
() =

Tangent – Sex marker discovery (preliminary) using Genome Wide Association Study



Scientific Reports 9, Article number: 116 (2019) Cite this article

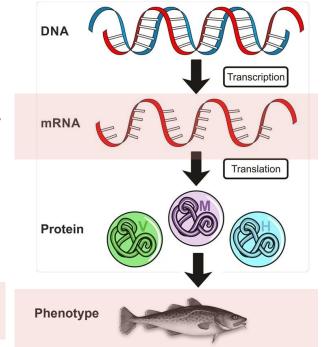
Benchtop sex assay in Atlantic cod



Pacific halibut assessment uses sex data based on DNA sample

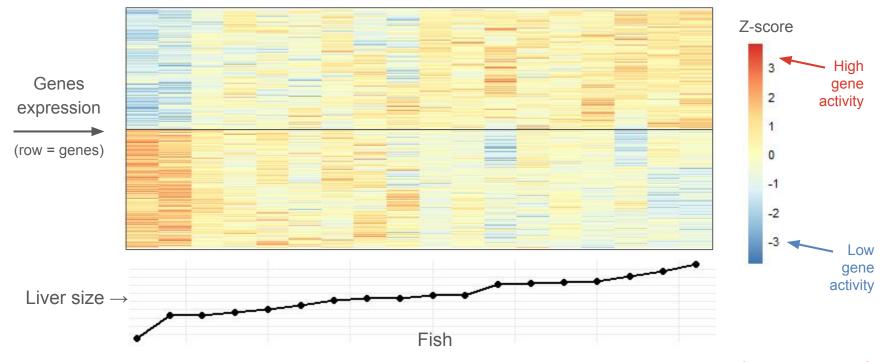
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~ 1,600 genes with <u>expression</u> associated with liver size in warming



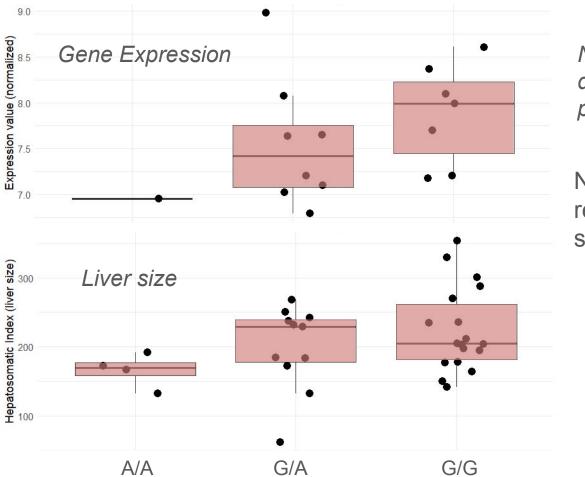
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Liver size performance indicators in warming, both GENETICS and EXPRESSION

Cell	Chromosome	# of markers	Gene ID	Protein Name	Function
adhesion	4	3	LOC132456135	Netrin receptor UNC5D	Cell adhesion, apoptosis in response to DNA damage
Calcium transport Immune system	12	2	tmco1	Calcium load-activated calcium channel	Calcium transport, endoplasmic reticulum calcium homeostasis
	10	1	LOC132466560	TBC1 domain family member 9B	Membrane trafficking, calcium transport
	2	1	LOC132453053	Stonustoxin subunit beta	May be related immune system function. From stonefish, toxic/fatal to mammals.
	5	1	LOC132457513	Stonustoxin subunit beta	May be related immune system function. From stonefish, toxic/fatal to mammals.
	23	1	LOC132452628	NLR family CARD domain-containing protein 3	Negative regulator of the innate immune response
	23	1	LOC132452644	HERV-H LTR-associating protein 2	Enhances T-cell proliferation and cytokine production
	17	2	LOC132445594	Unknown	Unknown

Liver size performance indicators in warming, both GENETICS and EXPRESSION

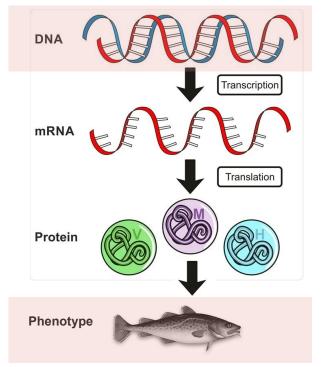
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NLR family CARD domain-containing protein 3

Negatively regulates immune system "Genome-to-Phenome" dataset for juvenile Pacific cod

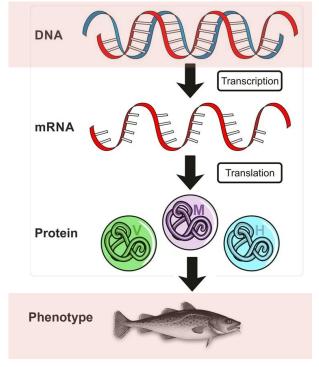
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- Gene expression Lipid usage, immune activity, & damage control may deplete energy reserves
- ✓ Markers of juvenile performance in warming
 - ~100 genetic markers
 - ✓ ~1,600 gene expression indicators



Adapted from: udaix/Shutterstock.com

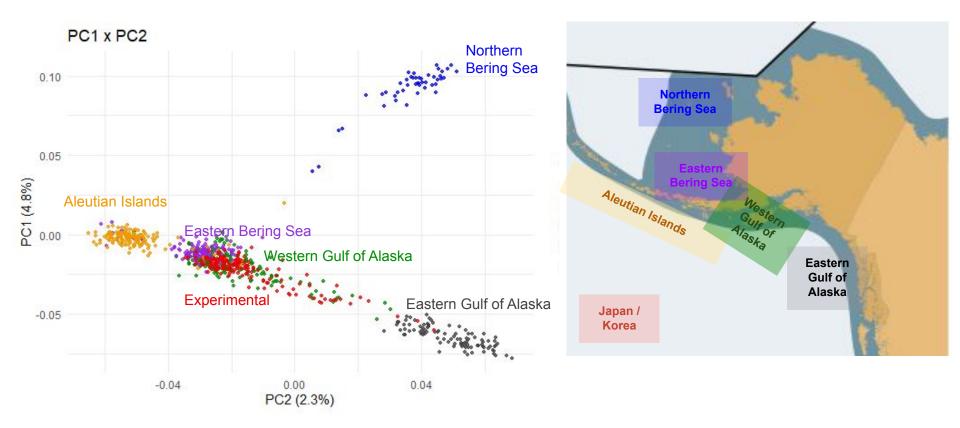
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- ✓ Markers of juvenile performance in warming
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 - ~1,600 gene expression indicators
- Can we predict "performance" or "resilience" of other cod groups using our markers?

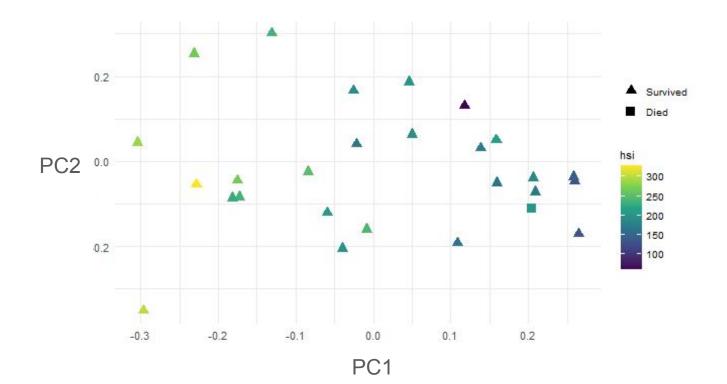


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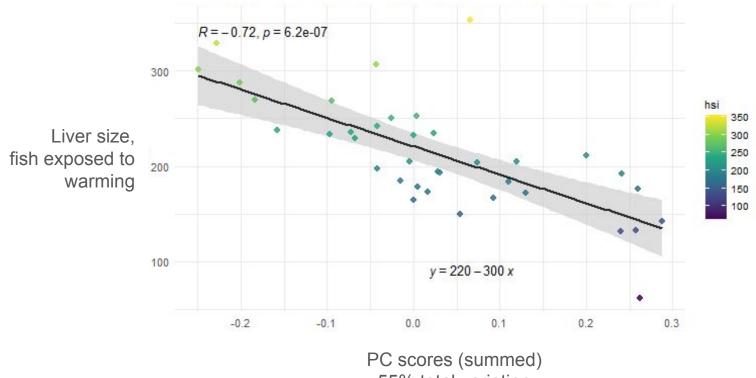
Which populations would we predict to have largest livers in warm conditions?



Analyses so far - use genotypes at **32 markers to predict liver size** with PC scores

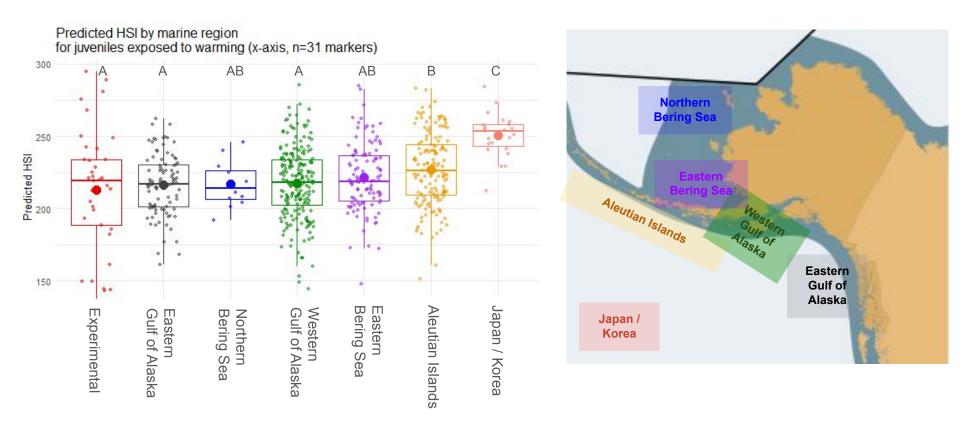


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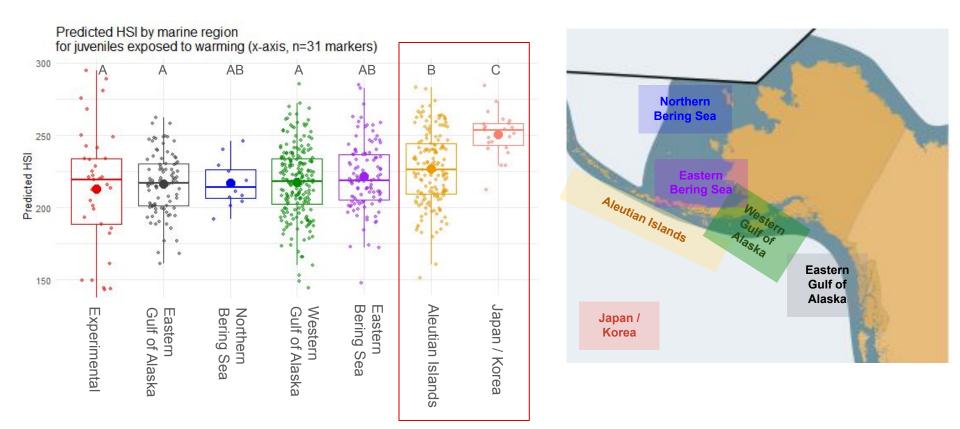


55% total variation

Which populations would we predict to have largest livers in warm conditions?



Which populations would we predict to have largest livers in warm conditions?





- Low juvenile recruitment in GoA during heatwaves is likely related to low overwintering survival due to fewer lipid reserves.
- Stock assessments may need to adjust mortality/recruitment rates in heat wave years
 - Larval study (Slesinger et al. 2024; Spencer et al. *in revision) <u>much</u> higher mortality*
- Genetic variability related to adaptive traits may enable selection for individuals more capable of allocating lipid reserves
- Other Pacific cod groups could be screened for **putative markers of performance**
 - Through time before/during/after heat waves (future project!) is selection happening?
 - Distinct Pacific cod groups are some groups more resilient than others?
- Building Resources for Pacific cod lists of genes that are temperature sensitive in larvae, juveniles, indicators of performance, etc.

Collaborators & Funders



Newport (Hatfield)

Ben Laurel Emily Slesinger Louise Copeman MaryBeth Hicks Tom Hurst

Seattle (Regional office)

Ingrid Spies Sara Schaal



University of Washington Steven Roberts





