Identification of molecular growth signatures in skeletal muscle of juvenile Pacific halibut (Hippoglossus stenolepis) for monitoring population growth patterns

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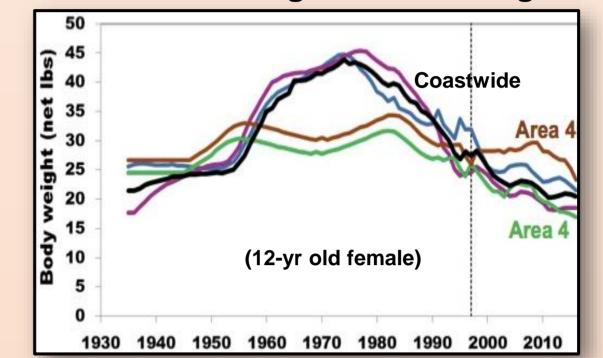
INTERNATIONAL PACIFIC



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Historical changes in size-at-age



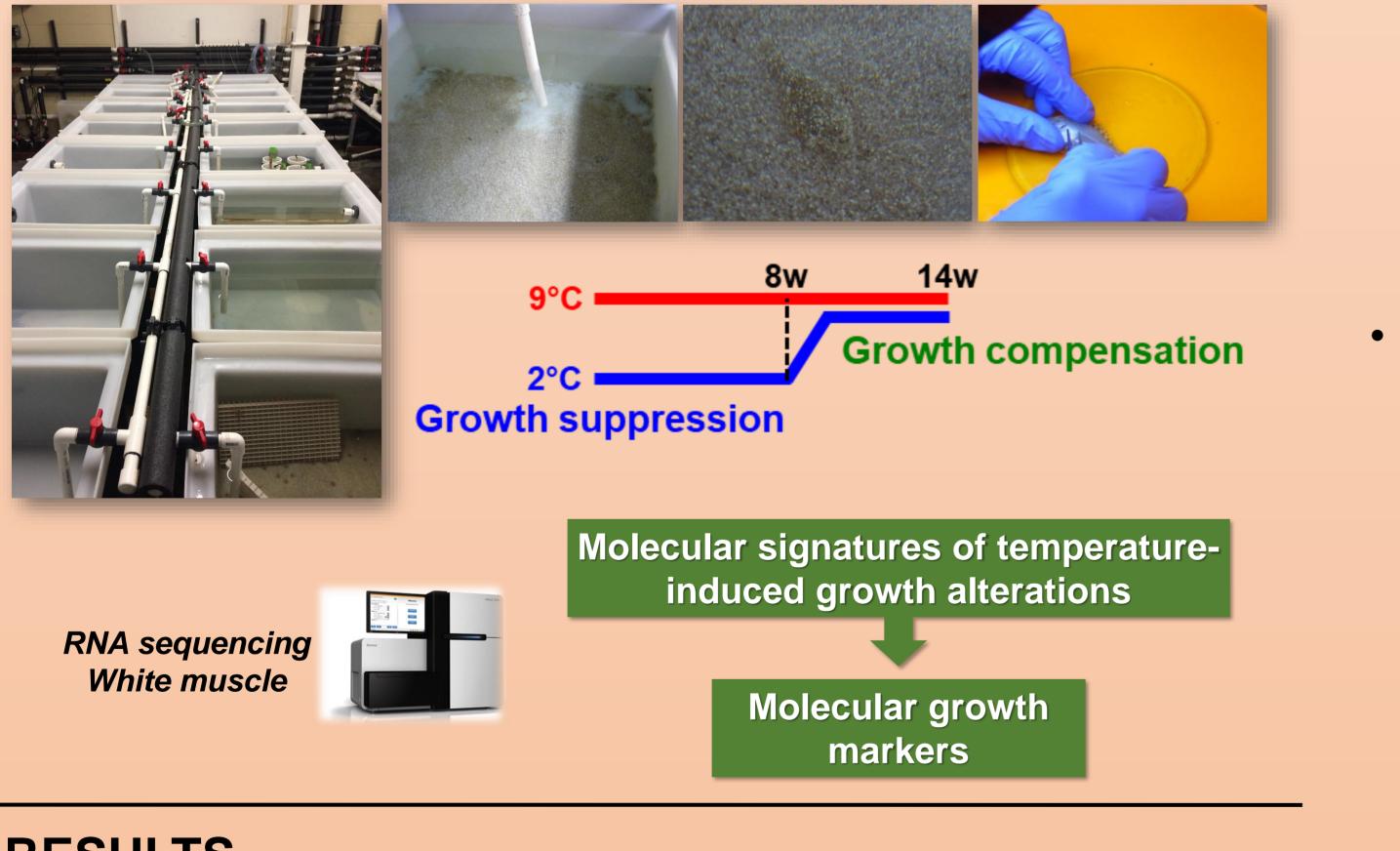


The International Pacific Halibut Commission has reported changes in the size-at-age (SAA) of Pacific halibut (*Hippoglossus*) stenolepis) caught in the commercial fishery as well as in its own survey research for almost 100 years. Although an increase in SAA was observed between the 1930's until the 1980's, SAA has significantly declined since the 1990's until today, as evidenced by a 50% reduction in body weight for a typical 12-year old female during this period (Figure 1). However, our understanding of the potential causes for the long-term variability in SAA is still rather scarce. Although a number of factors could be contributing to this variability, recent analyses have suggested that temperature variation may have been a contributing factor to the observed changes in SAA in the Pacific halibut. Therefore, there is an urgent need to better understand the physiological effects of temperature on growth in this species.

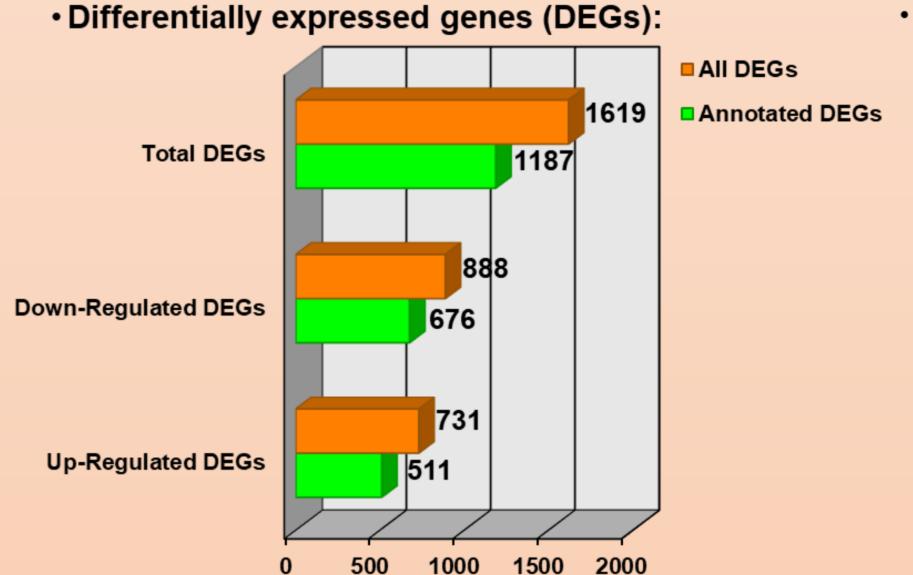
Figure 1. Coastwide changes in the average body weight of a 12-yr old female Pacific halibut (black line).

MATERIALS AND METHODS

Juvenile Pacific halibut of approximately 6 month of age were collected off the coast of Kodiak, Alaska, US and transferred to the aquatic facilities of the Hatfield Marine Science Center in Newport, Oregon, US. Individually pittagged fish were acclimated for 8 weeks to 2°C and 9°C in duplicate tanks (N = 5) prior to sampling. Subsequently, half of the fish previously acclimated at 2°C were gradually brought up to 9°C and held at 9°C for 6 additional weeks prior to sampling. The transcriptomic responses of white skeletal muscle from fish experiencing temperature-induced growth suppression and growth compensation were analyzed by RNA sequencing (Illumina).



Transcriptomic responses to temperature-induced growth suppression



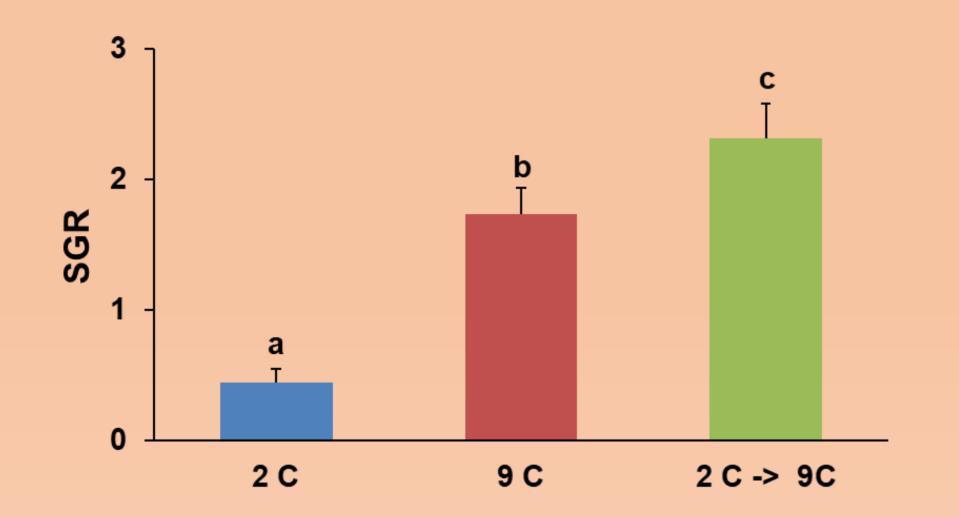
 Functional categories 	(Down-regulated at 2°C)
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Change	Functional category
Down	Muscle development and contraction
Down	Transcription and translation
Down	Protein and carbohydrate metabolism
Down	Energy metabolism and transfer
Down	Cell division
Down	Stress response
Down	Immune response

Figure 3. Left: Number of differentially expressed genes. Right: Functional categories

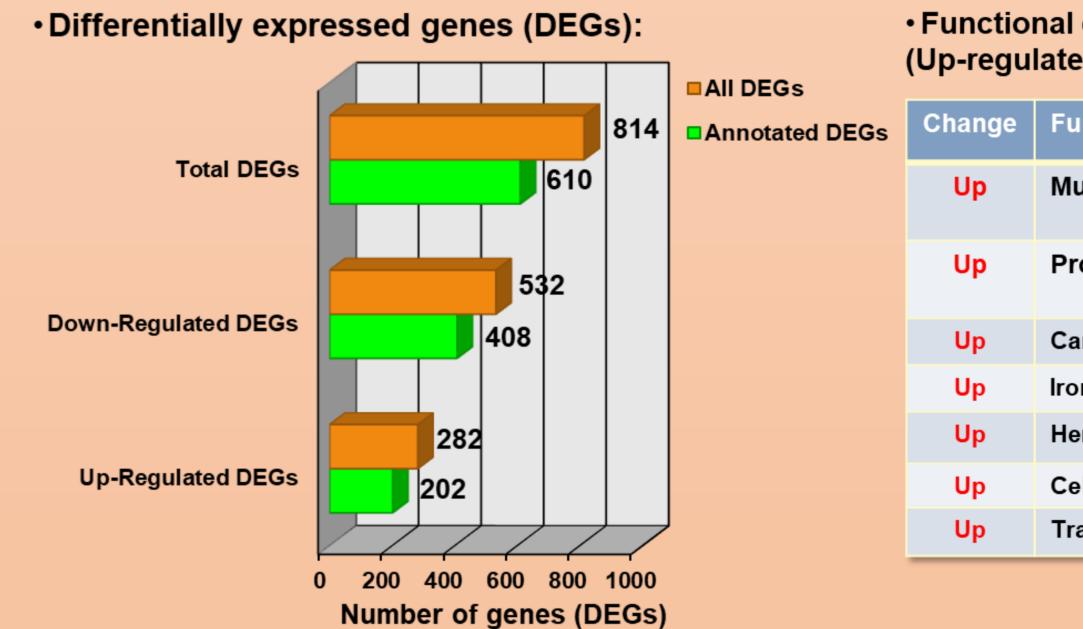
RESULTS

Temperature modulates the specific growth rate (SGR)



of genes significantly down-regulated under growth suppression.

Transcriptomic responses to temperature-induced growth stimulation



 Functional categories (Up-regulated from 2°C to 9°C):

Functional category Muscle development and contraction Protein metabolism and modification Carbohydrate metabolism (for ATP) Iron transport and binding Hemoglobin synthesis Cell adhesion and proliferation Transcription and translation

Figure 4. Left: Number of differentially expressed genes. Right: Functional categories of genes significantly up-regulated under growth stimulation.

CONCLUSIONS

• Acclimation at 2°C resulted in a significant reduction in the specific growth rate (SGR) whereas a significant increase in SGR was observed as a result of temperature-induced growth compensation.

Temperature

Figure 2. Effects of temperature on the specific growth rate in juvenile Pacific halibut. Different letters indicate statistically significant differences among the groups (N = 10).





- Growth suppression by low temperature acclimation is associated with a decrease in the expression of genes involved primarily in muscle function, protein synthesis, transcription and stress and immune response.
- Growth stimulation by temperature-induced compensation is associated with an increase in the expression of genes involved primarily in muscle structure and function and metabolic activation.
- The resulting molecular growth signatures will be useful to investigate potential changes in growth patterns in Pacific halibut.

ACKNOWLEDGEMENTS. This study was conducted with funding from IPHC and the North Pacific **Research Board (Project NPRB 1704).**