



Overview of the IPHC 5-year Biological and Ecosystem Science Research Program

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Joint NPFCM / IPHC Meeting

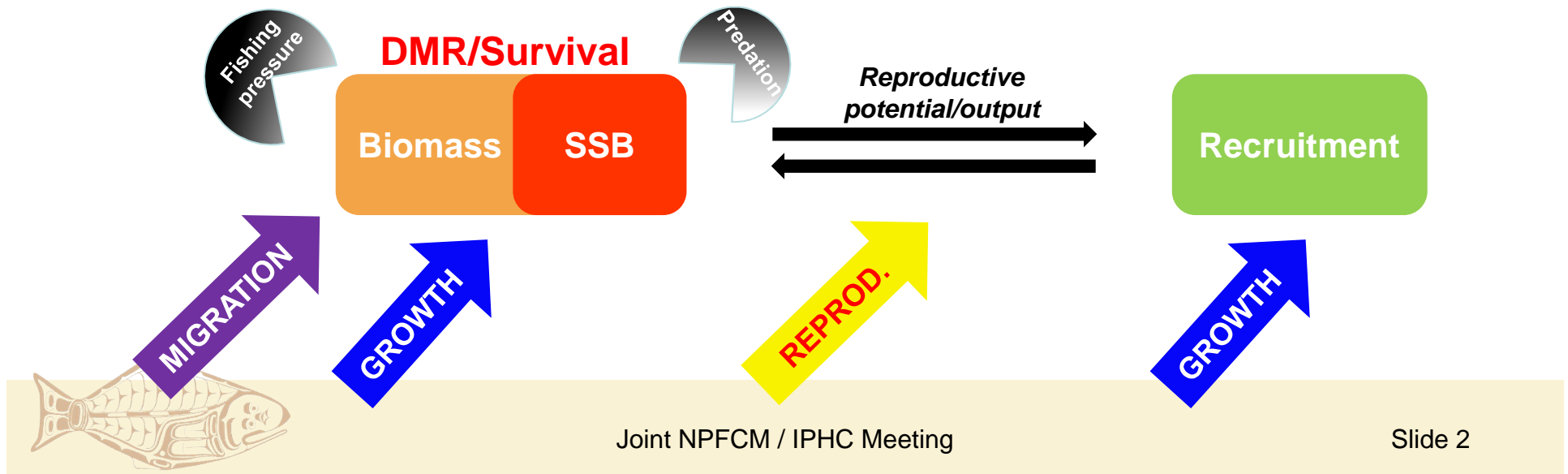
June 7, 2017

Primary research activities at IPHC

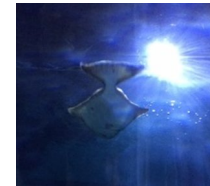


Primary objectives

- Identify and address *critical knowledge gaps* in the biology of the Pacific halibut
- Understand the influence of *environmental conditions* on halibut biology
- Apply resulting knowledge to reduce *uncertainty* in current stock assessment models



Primary research activities at IPHC



1. Reproduction

- SEX RATIO OF CATCH
- IMPROVED MATURATION ESTIMATES OF SPAWNING BIOMASS

2. Growth

- CHANGES IN SIZE AT AGE/BIOMASS
- TOOLS TO ASSESS FISH CONDITION

3. Migration

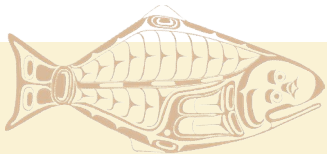
- LARVAL DISPERSAL
- ADULT FEEDING AND REPRODUCTIVE MIGRATION

4. DMRs and post-release survival assessment

- BYCATCH SURVIVAL ESTIMATES

5. Genetics and genomics

- GENETIC STRUCTURE OF THE POPULATION
- GENOMIC TOOLS (e.g. GENOME)



1. Reproduction

There are important knowledge gaps on the reproductive biology of the species

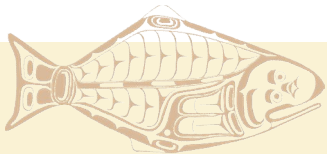
- SEX RATIO OF CATCH
- IMPROVED MATURATION ESTIMATES OF SPAWNING BIOMASS

What is needed?

- Knowledge on reproductive development, maturation, fecundity, sex determination mechanisms (sex identification), environmental and hormonal control of reproduction.
- Scientific-based criteria to identify reproductive status and potential.
- Updated estimates of age and size at maturation.
- Information on skipped spawning.

New proposed studies:

- *Full characterization of the annual reproductive cycle*
- *Identification of sex determination mechanism(s) and influencing factors*



1. Reproduction

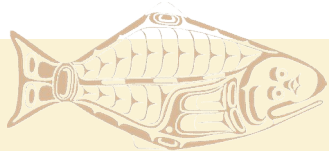
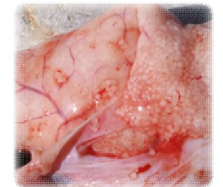
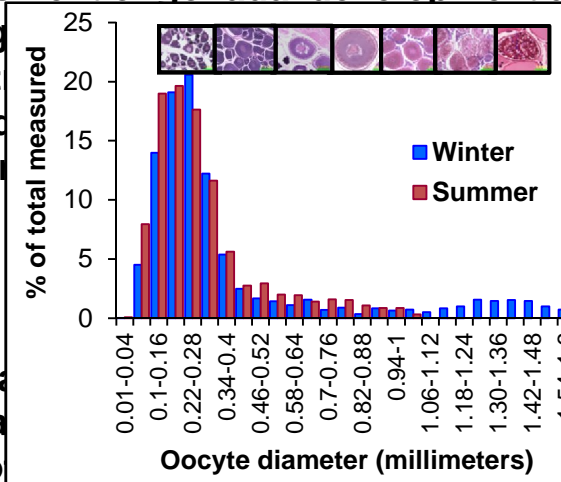
- **Full characterization of the annual reproductive cycle**

Objective: Understand temporal changes in reproductive development throughout an entire annual reproductive cycle in male and female Pacific halibut

- Histological assessment of gonadal development and maturation.
- Endocrine profiling
- Gene expression (t)
- Gonadosomatic index
- Ultrasound monitoring

Deliverables:

- Accurate spawning
- Updated management
- Estimates of
- Comprehensive reproductive monitoring of the adult population in order to improve our estimate of actual spawning biomass

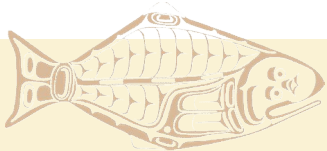
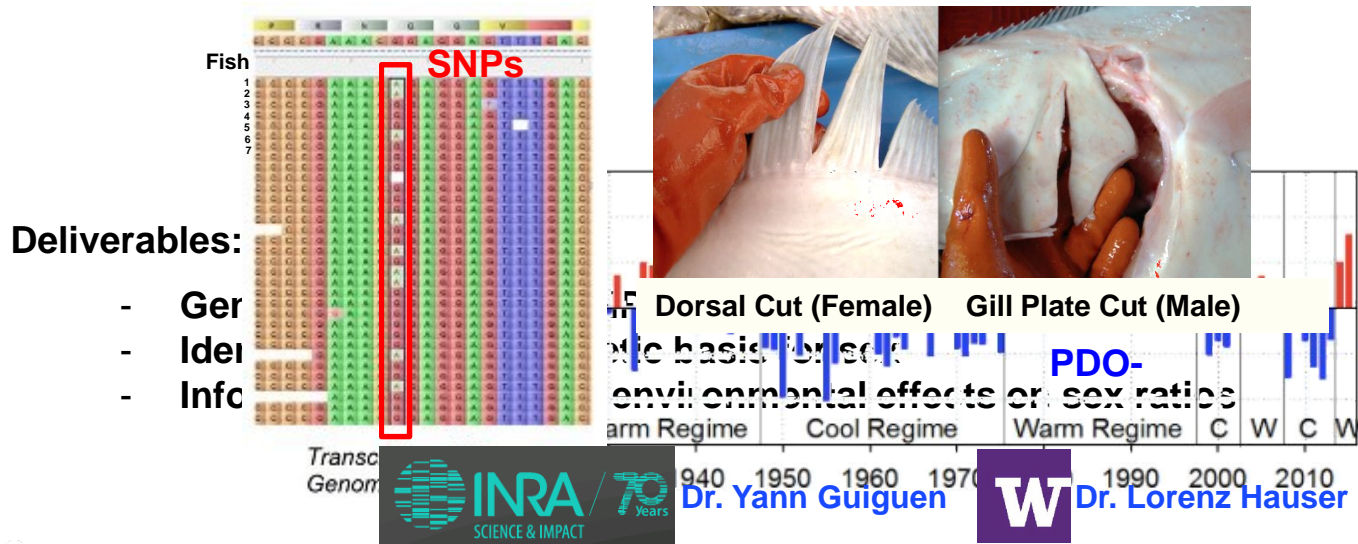


1. Reproduction

- **Identification of sex by genetic means and of sex determination mechanisms**

Objectives: To identify genetic markers for sex identification in commercial catch and to understand how sex is established in Pacific halibut

- Identification of genetic sex markers: Validation of the coast-wide sex-marking project



2. Growth

Little is known regarding what factors influence growth in this species

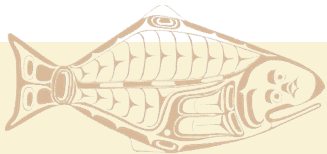
- CHANGES IN SIZE AT AGE/BIOMASS
- TOOLS TO ASSESS FISH CONDITION

What is needed?

- Knowledge on growth patterns and environmental influences.
- Improved understanding in the possible role of growth alterations in the observed decrease in size at age.

New proposed studies:

- *Extensive catalogue of physiological markers to monitor growth*
- *Evaluation of growth patterns and effects of environmental influences*



2. Growth: proposed studies

- **Catalogue of muscle growth studies.**

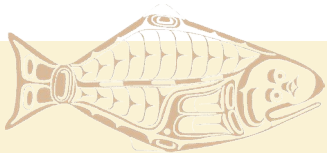
Objective: Identify

- Identification of growth-related genes
- Develop molecular assays

Annotation	Gene symbol	Length (nt)	Identity (%)	Function
Androgen receptor	<i>ar</i>	4426	81.48	Protein synthesis
Calcium/calmodulin-dependent protein kinase II alpha	<i>camk2a</i>	2342	87.27	Force transmission
Creatine kinase, muscle a	<i>ckma</i>	2256	89.76	Energy metabolism
Carnitine palmitoyltransferase 1B	<i>cpt1b</i>	762	87.81	Lipid metabolism
Dystrophin	<i>dmd</i>	1282	87.23	Force transmission
Eukaryotic translation initiation factor 4eb	<i>eif4eb</i>	1168	85.19	Protein synthesis
F-box protein 32	<i>fbxo32</i>	695	86.25	Protein atrophy
Glycogen synthase 1	<i>gys1</i>	3723	89.47	Energy metabolism
Histone deacetylase 1	<i>hdac1</i>	2490	96.35	Muscle repressor
Insulin-like growth factor 2 receptor	<i>igl2r</i>	511	100.00	Growth regulator
Insulin-like growth factor binding protein 5b	<i>igfbp5</i>	1372	81.5	Growth regulator
Lipoprotein lipase	<i>llpl</i>	1789	60.48	Lipid metabolism
Myocyte enhancer factor 2cb	<i>mef2cb</i>	564	79.8	Muscle growth
Myostatin b	<i>mstnb</i>	39	95.71	Growth regulator
Mechanistic target of rapamycin	<i>mtor</i>	1153	97.92	Protein synthesis
Myogenic factor 6	<i>myf6</i>	819	77.19	Muscle growth
Myosin, heavy polypeptide 1.3, skeletal muscle	<i>myh1.3</i>	246	86.42	Muscle growth
Myoblast determination protein 1 homolog	<i>myod</i>	2497	72.67	Muscle development
Myozenin 1a	<i>myoz1a</i>	766	74.6	Force transmission
Nuclear factor of activated T-cells, cytoplasmic 3	<i>nfatc3</i>	1587	62.96	Muscle activity
Paired box 3a	<i>pax3</i>	269	75	Muscle development
Paired box 7b	<i>pax7b</i>	297	85.71	Muscle development
Peroxisome proliferator-activated receptor gamma, coactivator 1 alpha	<i>ppargc1a</i>	519	88.7	Energy metabolism
Protein phosphatase 3, catalytic subunit, alpha isozyme	<i>ppp3ca</i>	3407	83.69	Muscle activity
Protein kinase, AMP-activated, alpha 1 catalytic subunit	<i>prkaa1</i>	1925	70.96	Energy metabolism
Phosphorylase, glycogen, muscle	<i>pygma</i>	5514	90.91	Energy metabolism
Serum response factor	<i>srf</i>	4393	63.81	Muscle development
Transforming growth factor, beta 1a	<i>tgfb1a</i>	561	77.04	Growth regulator
Tripartite motif containing 63b	<i>trim63b</i>	2117	81.16	Protein atrophy

Deliverables:

- Establishment of a growth-related gene sequence dataset
- Molecular assays to monitor growth patterns based on growth-markers



2. Growth

- **Evaluation of growth patterns and effects of environmental influences**

Objective: Identify molecular, biochemical and isotopic profiles characteristic of specific growth patterns and evaluate potential effects of environmental influences.

- Evaluation of different growth patterns in the *wild*.

In BS NMFS trawl survey in 2016:

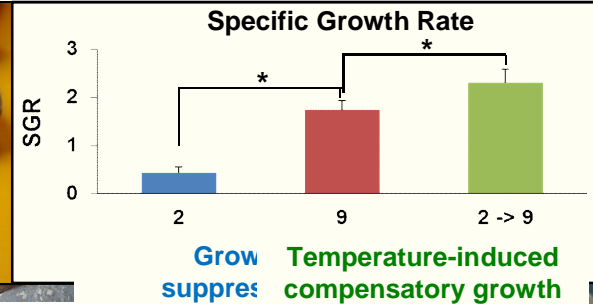
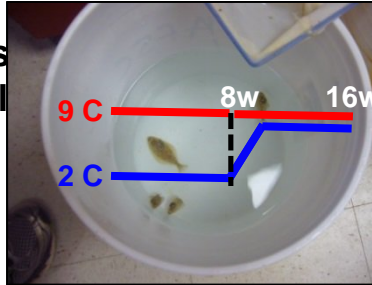
- - 75 fish <40 cm length
- - 75 fish 40-60 cm length
- - 75 fish 60-80 cm length

Continued in 2017



Characterization of molecular and biochemical growth markers in muscle samples from age-matched individuals

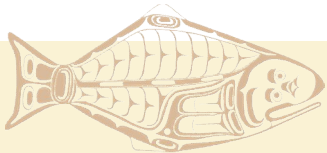
- Establish molecular



Grow suppress Temperature-induced compensatory growth

- Isotopic tissue turnover to trace dietary and/or habitat shifts

^{13}C , ^{15}N



2. Growth

- Investigate the effects of **environmental factors** on growth performance.
 - Effects of **temperature, salinity, dissolved oxygen and water pH** on growth.



- *Identify the optimal environmental conditions for growth.*
- Understand the basis of **sexual dimorphic growth** in the Pacific halibut.

Male



Female



Deliverables:

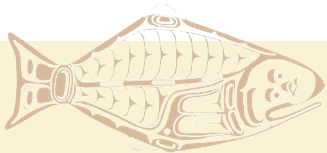
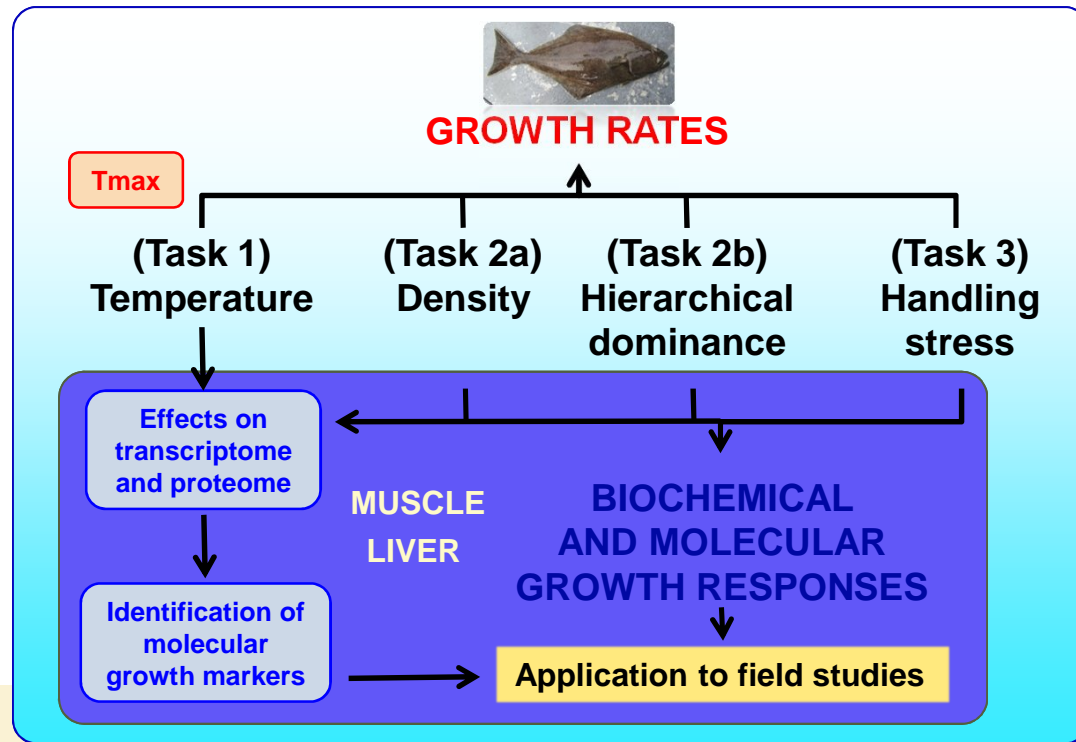
- Identification and validation of growth markers for field studies
- Characterization of molecular and biochemical growth signatures
- Environmental effects on somatic growth
- Improved biological inputs on biomass estimates



2. Growth



- NPRB Funding (2017-2019):** “Somatic growth processes in the Pacific halibut (*Hippoglossus stenolepis*) and their response to temperature, density and stress manipulation effects”. IPHC (coordinator) / AFSC – Newport, OR



3. Migration

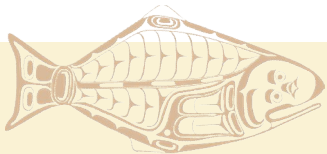
- LARVAL DISPERSAL
- ADULT FEEDING AND REPRODUCTIVE MIGRATION

What is needed?

- Improve our understanding on larval, juvenile and reproductive migration.
- Incorporate additional sources of biological information on migration studies.

Projects:

- *Juvenile and adult feeding migrations*
- *Reproductive and annual migrations of adult fish*
- *Larval migration and connectivity*



3. Migration

- **Juvenile and adult migration studies**

- Juvenile wire tagging: • NMFS trawl tagging project
- Adult wire tagging: • IPHC survey tagging project
 - 2016 pilot study in area 4D (U32)
 - 2017 coast-wide study (U32)

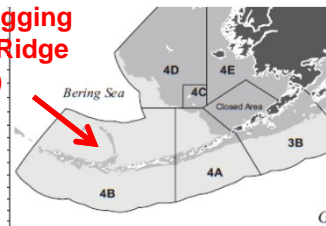
Fin clips are collected: Genetic analyses of tagged fish to shed light on migration patterns and geographic origin.



- Tail pattern recognition



- PAT tagging Bower's Ridge (4B)



- **Reproductive and annual migration**

Deliverables:

- Improved knowledge on juvenile, adult and reproductive migrations and identification of spawning areas

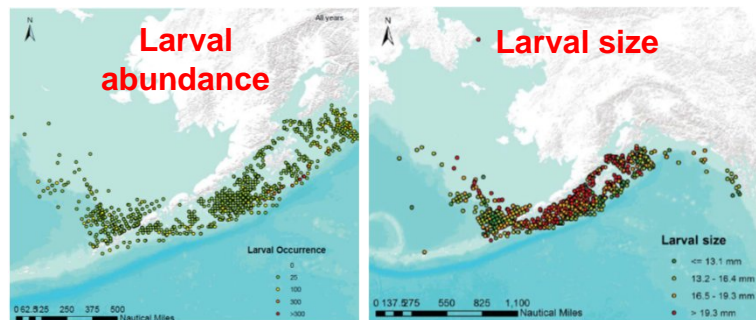


3. Migration

- *Larval migration and connectivity*

Objective: Understand the mechanisms of larval connectivity between GOA and BS.

- Collect data from the NMFS ichthyoplankton survey and map larval distribution over time and space.
- Collect larval samples from the survey to conduct genetic analyses.



Collaboration with Janet Duffy-Anderson, Esther Goldstein, William Stockhausen (NOAA-AFSC-Seattle)

Deliverables:

- Improved knowledge on larval distribution, migration and genetic structure within the population



4. DMRs and survival assessment

Little is known regarding the factors that influence bycatch survival

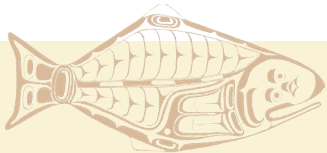
• BYCATCH SURVIVAL ESTIMATES

What is needed?

- To introduce quantitative measurable factors that are linked to fish handling practices and to fish physiological condition and ultimately to survival in order to improve current DMR estimations

Projects:

- *Evaluation of the effects of fish handling practices on injury levels and the physiological condition of captured Pacific halibut*
- *Investigate the relationship between physiological condition post-capture and survival as assessed by the use of accelerometer tags.*
- **FUTURE: Improving estimates of survival of fish caught in the trawl fishery**



4. DMRs and survival assessment

- *Evaluation of the effects of fish handling practices on injury levels and the physiological condition of captured Pacific halibut*

Objective: Understand relationship between handling practices and physiological condition of captured Pacific halibut in the longline fishery

- Assess injuries associated with release techniques (gangion cut, careful shake, hook straightening).
- Determine the physiological condition of all captured fish with associated injury levels after different deck exposure times: condition factor index (Kn), energy (fat) levels, morphometric analyses.
- Measure the levels of stress and physiological disturbance indicators in the blood of all captured fish (cortisol, lactate, glucose, potassium, hematocrit).

Deliverables:

- Injury profile for different release techniques in the longline fishery
- Physiological assessment of fish handling practices: **fish condition index post-capture**



4. DMRs and survival assessment

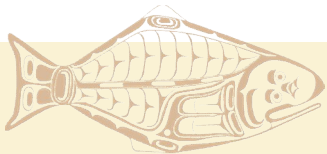
- ***Investigate the relationship between physiological condition post-capture and survival as assessed by tagging.***

Objective: Measure post-release survival in Pacific halibut and relate it to physiological condition and capture-related events

- Tag fish that have been exposed to different handling practices in the longline fishery with accelerometer tags in addition to conventional tags (wire).
- Assess survival of fish according to size and physiological condition.

Deliverables:

- Information on post-release survival in relation to handling practices and physiological condition.
- Information on post-release survival in relation to size.
- Estimating DMRs by EM.

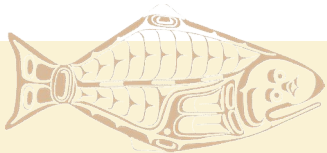
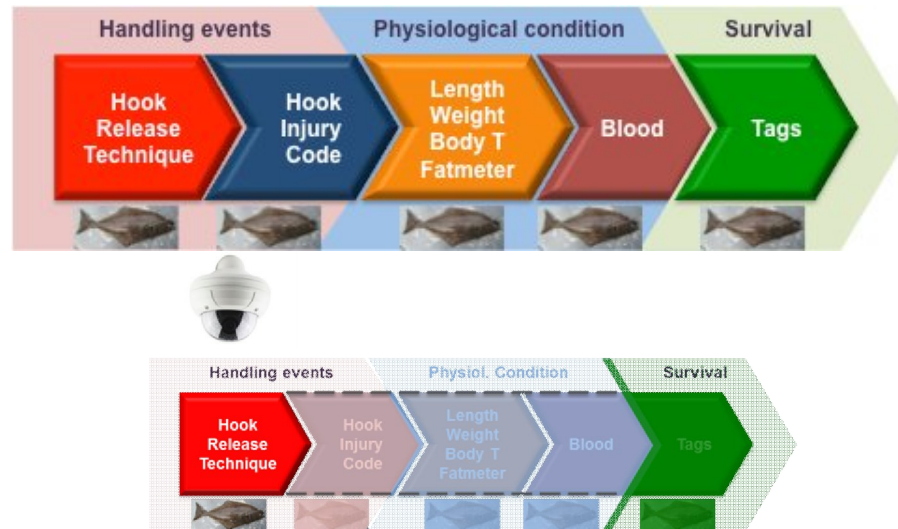


3. DMRs and survival assessment



- **Saltonstall – Kennedy Program Funding (2017-2019):** “Improving discard mortality rate estimates in the Pacific halibut by integrating handling practices, physiological condition and post-release survival”. IPHC (coordinator) / APU – Anchorage, AK

Dr. Brad Harris
Dr. Nathan Wolf



3. DMRs and survival assessment

- *Improving estimates of survival of Pacific halibut in the trawl fishery*

Objective: Assess the condition of discarded Pacific halibut in the non-directed trawl fishery and

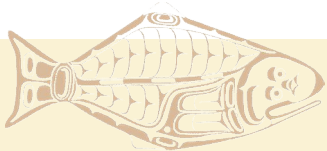


discarded Pacific halibut in the estimates of discard mortality rates

- Continue and capitalize on collaborative research on (e.g. Amendment 80 fleet) to plan
- Apply methods to assess captured halibut.
- Determine survival rates of discarded halibut after tagging.
- Relate physiological condition with survival rates of discarded halibut

Deliverables:

- Improved knowledge of survival of discarded halibut and, consequently, improved estimates of discard mortality rates in the trawl fishery.



5. Genetics and genomics

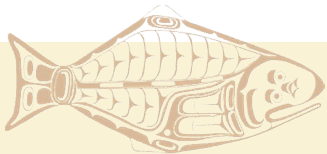
- GENETIC STRUCTURE OF THE POPULATION
- GENOMIC TOOLS (e.g. GENOME)

What is needed?

- Improved knowledge on the genetic composition of the population
- Establish genomic resources for the species
- Genome-wide association studies to evaluate genetic effects of fishery-dependent and fishery-independent influences on growth, reproduction, nutrition, etc.

New proposed studies:

- *Population genetic studies*
- *Sequencing of the Pacific halibut genome*

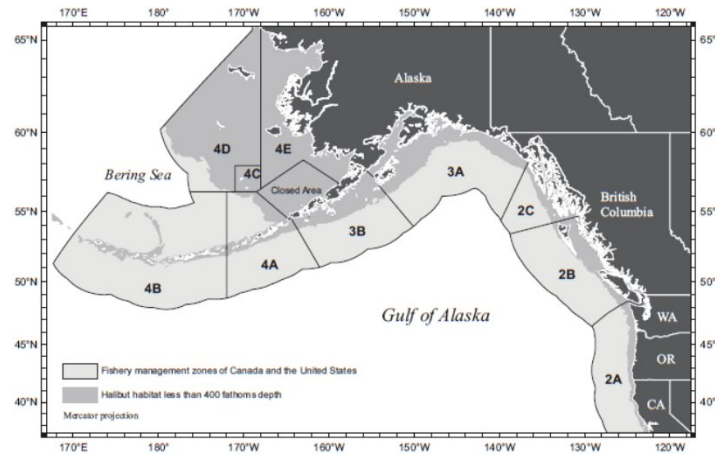


5. Genetics and genomics

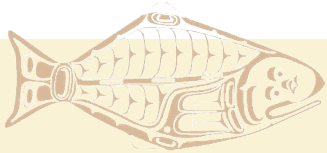
- *Population genetic studies*

Objective: Genetic characterization of Pacific halibut throughout its distribution range

- Characterization of population structure by RAD sequencing and SNP analysis.
- Identification of possible genetic signatures of geographical origin



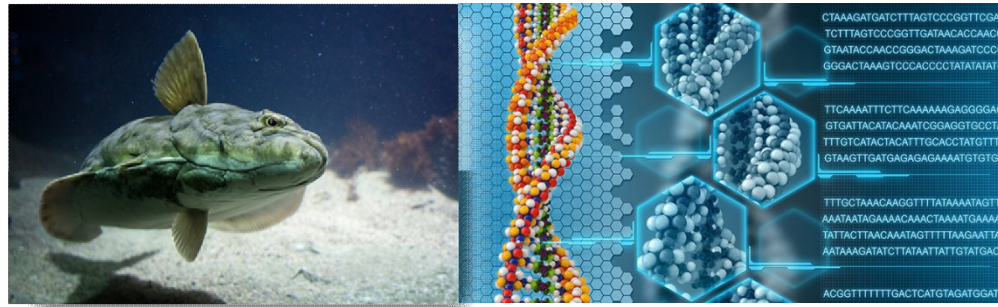
Dr. Lorenz Hauser



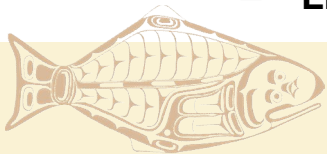
5. Genetics and genomics

- *Pacific halibut genome*

Objective: Obtain a first draft sequence of the Pacific halibut genome



- Provide genomic resolution to genetic markers (SNPs or transcripts).
- Identify genomic regions and genes responsible for temporal and spatial adaptive characteristics.
- Genome-wide association studies to try to understand the genetic basis of growth, reproductive performance, migratory behavior and performance, etc.
- Link genotype and phenotype.

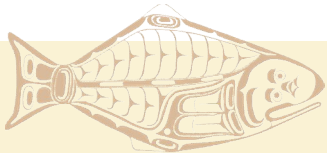
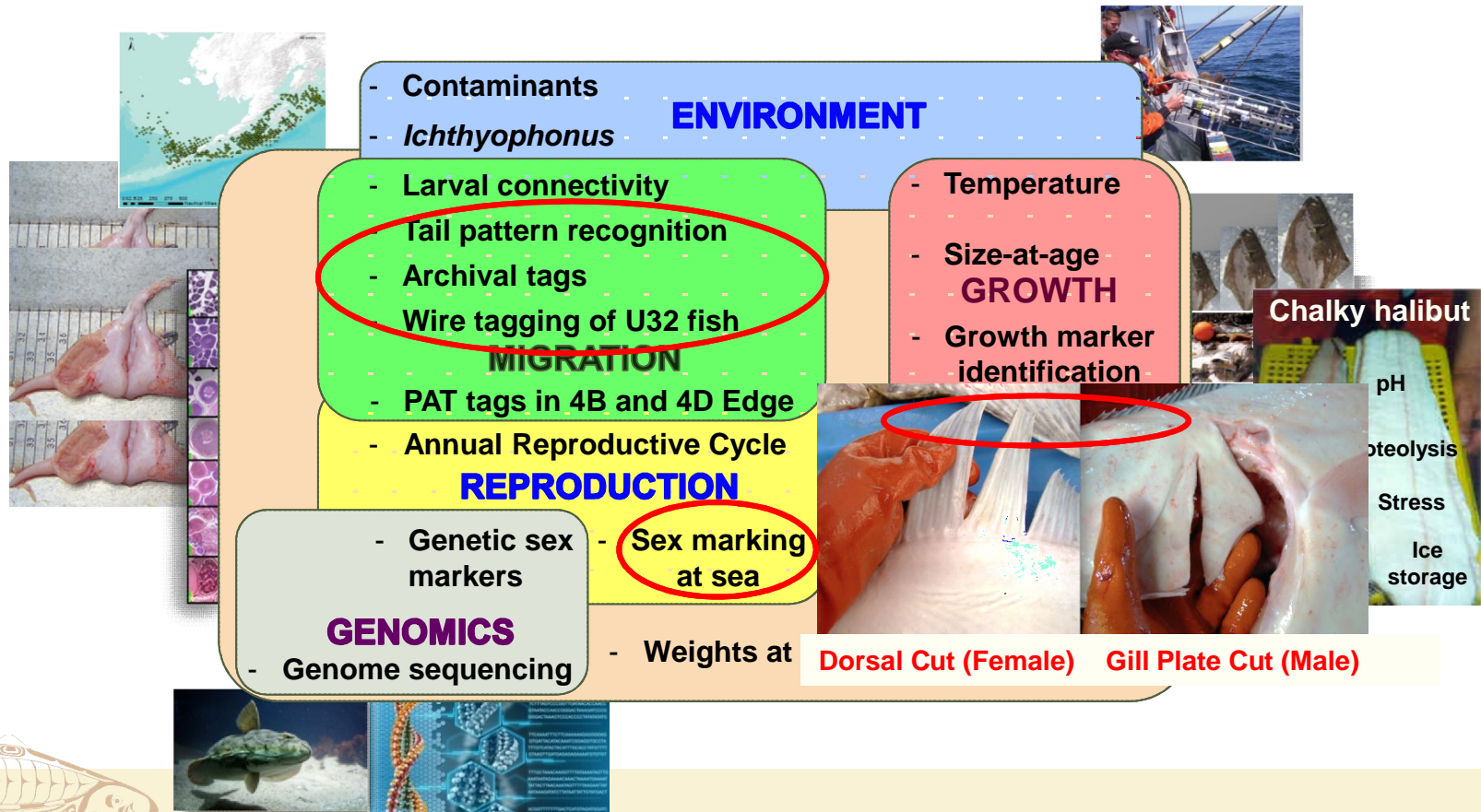


Dr. Yann Guiguen



Dr. Lorenz Hauser

Research projects for 2017



Temporal chart of activities

	2016	2017	2018	2019	2020	2021
Reproduction		Annual reproductive cycle				
		Sex determination mechanisms				
	Sex identification					

