



5th International Conference on Integrative Salmonid Biology
Monday 11 March – Thursday 14 March 2024
Bell Harbor International Conference Center

Monday 11 March		Location
14:00 – 19:00	Registration Open	Pre-Function Lobby
17:30 – 19:30	Meet and Greet	Pre-Function and Sound

Tuesday 12 March		Location
07:30 – 19:30	Registration Open	Pre-Function Lobby
07:30 - 08:30	Breakfast	Harbor
08:30 – 09:00	<p>Welcome and Introduction: Kerry Naish and Krista Nichols</p> <p>Welcome by Programme Sponsors: Tony Brooks, Genome British Columbia</p> <p>Conference Background and History: Ben Koop, University of Victoria</p>	Bay Auditorium
09:00 – 10:20	<p>Session One: Genome resources, functional genomics, and evolution of salmonids</p> <p><i>Moderator: Ben Koop</i></p>	Bay Auditorium
09:00 – 09:20	<p>T-Invited-1 Towards complete chromosome sequences and pan-genomic resources for salmonids</p> <p><i>Sigbjørn Lien, Norwegian University of Life Sciences, Norway</i></p>	
09:20 – 09:40	<p>T-Invited-2 Genomic analysis through the lens of pangenome</p> <p><i>Maria Simak, Universidad de Chile, Chile</i></p>	
09:40 – 10:00	<p>T-Invited-3 Pan-genomics reveals hidden genomic variation in Atlantic salmon</p> <p><i>Kristina Stenløkk, Norwegian University of Life Sciences, Norway</i></p>	
10:00 – 10:20	<p>T-Invited-4 Functional annotation of salmonid genomes: a window into regulatory changes following autopolyploidization</p> <p><i>Dan Macqueen, Roslin Institute, University of Edinburgh, UK</i></p>	



10:20 – 10:50	Networking/Exhibition/Refreshments	
10:50 – 12:05	Session Two: Genome resources, functional genomics, and evolution of salmonids <i>Moderator: Sigbjørn Lien</i>	Bay Auditorium
10:50 – 11:10	T-Invited-5 New rainbow trout genome assembly allows tracing 100 million years of gene evolution following whole genome duplication <i>Ali Ali, University of Maryland, USA</i>	
11:10 – 11:30	T-Invited-6 Wound atlas of the skin of Atlantic salmon <i>Rose Ruiz Daniels, Institute of Aquaculture, University of Stirling, UK</i>	
11:30 – 11:50	T-Invited-7 Uses of biotechnology to control reproduction in Atlantic salmon <i>Lene Kleppe, Institute of Marine Research, Norway</i>	
11:50 – 12:05	T-Oral-1 Single-cell transcriptomics indicates lasting impact of embryonic temperature on salmonid health <i>Carl Milton, University of Edinburgh, UK</i>	
12:20 – 13:20	Networking/Exhibition/Lunch	
13:20 – 14:55	Session Three: Functional genomics and annotation/Advances in aquaculture <i>Moderators: Dan Macqueen</i>	Bay Auditorium
13:20 – 13:35	T-Oral-3.1 The epigenome changes during interferon response genes in Atlantic salmon <i>Shahmir Naseer, University of Aberdeen, UK</i>	
13:35 – 13:50	T-Oral-3.2 Development of full-length nanopore RNA-Seq for alternative isoform discovery in Atlantic salmon <i>Oliver Eve, University of Edinburgh, UK</i>	
13:55 – 14:15	T-Invited-8 Implementing new technologies into existing structures – how, when and why? <i>Serap Gonen, Aquagen, Norway</i>	



14:15 – 14:35	T-Invited-9 Beyond genomic selection: Next steps in aquaculture breeding <i>José Manuel Yáñez, Universidad de Chile, Chile</i>	
14:35 – 14:55	T-Invited-10 Improving production efficiency and thermotolerance in New Zealand Chinook salmon <i>Jane E Symonds, Cawthorn Institute, New Zealand</i>	
14:55 – 15:25	Networking/Exhibition/Refreshments	
15:25 – 16:50	Session Four: Advances in aquaculture <i>Moderators: José Manuel Yáñez</i>	Bay Auditorium
15:25 – 15:45	T-Invited-11 Mechanisms of interspecific variation in sea lice resistance revealed by single-cell transcriptomics <i>Sarah Salisbury, Roslin Institute, University of Edinburgh, UK</i>	
15:45 – 16:05	T-Invited-12 Leveraging microbiota services through hologenomic analyses in farmed fish <i>Morten Limborg, University of Copenhagen, Denmark</i>	
16:05 – 16:20	T-Oral-6 Loss of FSHR prevents maturation in male Atlantic salmon <i>Ana Wargelius, Institute of Marine Research, Norway</i>	
16:20 – 16:35	T-Oral-5.4 Lack of VGLL3a delays onset of male maturation in Atlantic salmon <i>Erik Kjærner-semb, Institute of Marine Research, Norway</i>	
17:00 – 19:30	Poster Networking Session, Welcome Reception	Sound and Pre-Function Lobby

Wednesday 13 March		Location
08:00 – 17:00	Registration Open	Pre-Function Lobby



07:30 - 08:45	Breakfast	Harbor
08:30– 10:10	Session Five: Environment, Ecology and Conservation <i>Moderators: Kerry Naish and Krista Nichols</i>	Bay Auditorium
08:30 – 08:40	SESSION DEDICATION – Prof. Louis Bernatchez	
08:40 – 09:10	Keynote Session – Robin Waples, <i>The salmon genomics revolution: Prediction vs reality</i>	
09:10 – 09:30	W-Invited-13 Sustainable development of small-scale commercial indigenous Arctic fisheries <i>Stephan Schott, Carleton University, Canada</i>	
09:30 – 09:50	W-Invited-14 Complementary insights from genomics and Indigenous knowledge to support salmonid fisheries stewardship in northern Canada <i>Dylan Fraser, Concordia University, Canada</i>	
10:10 – 10:40	Networking/Exhibition/Refreshments	
10:40 – 12:10	Session Six: Environment, ecology and conservation <i>Moderators: Kerry Naish and Krista Nichols</i>	Bay Auditorium
10:40 – 11:00	W-Invited-15 Limiting river access to escapees: Consequences for an admixed Atlantic salmon population <i>Marine Brieuc, Institute of Marine Sciences, Norway</i>	
11:00 – 11:20	W-Invited-16 The good, the bad & the ugly of non-native salmonids in South America <i>Daniel Gomez-Uchida, Universidad de Concepción, Chile</i>	
11:20 – 11:40	W-Invited-17 Insights into run timing diversity revealed by range-wide Chinook salmon whole genomes <i>Tasha Thompson, Wild Salmon Center, USA</i>	
11:40 – 11:55	W-Invited-18 Beyond GWAS: Complex mechanisms underlie the simple genomic architecture of Atlantic salmon age at maturity <i>Craig Primmer, University of Helsinki, Finland</i>	
11:55 – 12:10	W-Oral-8 Evidence of genetic adaptation in <i>Oncorhynchus kisutch</i> exposed to 2013-2015 marine heatwave <i>Michelle Tsz Ting Chan, Simon Fraser University, Canada</i>	



12:10 – 13:10	Networking/Exhibition/Lunch	
13:10 – 14:45	Session Seven: Environment, Ecology and Conservation <i>Moderators: Kerry Naish and Krista Nichols</i>	Bay Auditorium
13:10 – 13:30	W-Invited-19 Documenting indigenous salmon stewardship practices through archaeology, ancient genomics and traditional knowledge <i>Camilla Speller, University of British Columbia, Canada</i>	
13:30 – 13:45	W-Oral-9 Genetic variation associated with adult migration timing in lineages of Steelhead & Chinook salmon in the Columbia River <i>Shawn Narum, Columbia River Inter-Tribal Fish Commission, USA</i>	
13:45 – 14:00	W-Oral-10 The distribution of adaptive genomic variation in space and time <i>Devon Pearse, NOAA, National Marine Fisheries Service, Southwest Fisheries Science Center, USA</i>	
14:00 – 14:15	W-Oral-11 Genomic basis of Canadian Atlantic salmon migration: Implications for climate-driven adaptation <i>Samantha Beck, UHI Inverness, UK and Department of Fisheries and Oceans/Dalhousie University, Canada</i>	
14:15 – 14:30	W-Oral-12 Genomic evidence for domestication selection in three hatchery populations of Chinook salmon <i>Charlie Waters, NOAA, National Marine Fisheries Service, Alaska Fisheries Science Center, USA</i>	
14:30 – 14:45	W-Oral-13 Major locus impacting run timing is conserved between Pink and Sockeye salmon <i>Natasha Howe, Pacific States Marine Fisheries Commission, USA</i>	
14:45 – 15:15	Networking/Exhibition/Refreshments	
15:15 – 16:15	Session Eight: Environment, ecology and conservation <i>Moderators: Kerry Naish and Krista Nichols</i>	
15:15 – 15:25	W-Flash-7 Evolutionary history and population structure of Fraser River salmon revealed by resequencing <i>Kris Christensen, University of Victoria, Canada</i>	
15:25 – 15:35	W-Flash-9 Genomic divergence of hatchery- and natural-origin Chinook salmon in two supplemented populations	



	<i>Michael Ford, NOAA, National Marine Fisheries Service, Northwest Fisheries Science Center, USA</i>
15:35 – 15:45	W-Flash-11 DNA methylation variation in hatchery Coho salmon and its potential heritability <i>Kyle Wellband, Department of Fisheries and Oceans, Canada</i>
15:45 – 15:55	W-Flash-13 Eco-evolutionary dynamics of dispersal: Insights from molecular pedigrees of wild Sockeye metapopulations <i>Samuel May, University of Washington/USDA Agricultural Research Service, USA</i>
15:55 – 16:05	W-Flash-14 Microbial community dynamics in a salmonid enhancement hatchery system <i>Shelby Reimer, BC Centre for Aquatic Health Sciences, Canada</i>

Thursday 14 March		Location
08:00 – 17:00	Registration Open	Pre-function Lobby
07:30 - 08:30	Breakfast	Harbor
08:30 – 10:05	Session Nine: Physiology and nutrition <i>Moderator: Sam Martin</i>	Bay Auditorium
8:30 – 8:50	TH-Invited-20 Gametes of semelparous Pacific salmon are repeatedly produced by surrogate Rainbow Trout <i>Goro Yoshizaki, Tokyo University of Marine Science and Technology, Japan</i>	
8:50 – 9:10	TH-Invited-21 Roles of miRNAs in reproduction <i>Julien Bobe, INRAE, France</i>	
9:10 – 9:30	TH-Invited-22 Cardiac morphological remodeling by environmental factors in Atlantic salmon <i>Ida Beitnes Johansen, Norwegian University of Life Sciences, Norway</i>	



9:30 – 9:50	TH-Invited-23 Chaperone-mediated autophagy regulates metabolism, energy balance, and oxidative stress in Rainbow Trout <i>Beth Cleveland, USDA, National Center for Cool and Coldwater Aquaculture, USA</i>	
9:50 – 10:05	TH-Oral-14 Impact of nutritional programming on intestinal transcriptome of Atlantic salmon (<i>Salmo salar</i>) <i>Marwa Tawfik, University of Aberdeen, UK</i>	
10:05 – 10:35	Networking/Exhibition/Refreshments	Harbor
10:35 – 12:05	Session Ten: Physiology and nutrition / Immunology, disease and host-pathogen interactions <i>Moderators: Yniv Palti</i>	Bay Auditorium
10:35 – 10:50	TH-Oral-15 Consequences of the embryonic environment in Atlantic salmon <i>Erik Burgerhout, Norwegian Institute of Food, Fisheries and Aquaculture Research, Norway</i>	
10:50 – 11:05	TH-Oral-16 Employing CRISPR genetic screens to identify mechanisms of environmental resilience in salmonids <i>Michael Phelps, Washington State University, USA</i>	
11:05 – 11:15	TH-Flash-15 Characteristics of gill tissue regeneration in Atlantic salmon (<i>Salmo salar</i>) <i>Ensiyeh Ghanizadeh-Kazerouni, The University of British Columbia, Canada</i>	
11:15 – 11:25	TH-Flash-16 Nutritional solutions to thermal stress in Atlantic salmon <i>Nick Wade, The University Of Edinburgh, UK</i>	
11:25 – 11:45	TH-Invited-24 Moving sterile Crispr salmon from tank to sea cage environment - regulatory barriers and social impacts <i>Anna Wargelius, Institute of Marine Research, Norway</i>	
11:45 – 12:05	TH-Invited-25 What have we learned about T Lymphocytes in salmonid fishes? <i>Monica Imarai, University of Santiago Chile, Chile</i>	
12:05 – 13:05	Networking/Exhibition/Lunch	Harbor
13:05 – 14:35	Session Eleven: Immunology, disease and host-pathogen interactions	Bay Auditorium



<i>Moderators: Beth Cleveland</i>	
13:05 – 13:25	<p>TH-Invited-26 Host responses associated with the evolution of viral generalism and specialism</p> <p><i>David Paez, US Geologic Survey/University of Washington, USA</i></p>
13:25 – 13:45	<p>TH-Invited-27 How can single cell genomics help us understand the pathogenesis of infectious salmon anaemia and other viral infections?</p> <p><i>Johanna Hol Fosse, Norwegian Veterinary Institute, Norway</i></p>
13:45 – 14:05	<p>TH-Invited-28 Genomic regulatory changes underlying the proinflammatory immune response in Atlantic salmon.</p> <p><i>Sam Martin, University of Aberdeen, UK</i></p>
14:05 – 14:20	<p>TH-Oral-17 Exploring Atlantic salmon head kidney cellular diversity by single-cell and single-nucleus transcriptomics.</p> <p><i>Adriana Magalhães Santos Andresen, Veterinærinstituttet, Norway</i></p>
14:20 – 14:35	<p>TH-Oral-18 Atlantic salmon spleen immune cell heterogeneity and cell-specific responses to bacterial infection revealed by single-nucleus transcriptomics</p> <p><i>Jianxuan Sun, and the Royal (Dick) School of Veterinary Studies, The University of Edinburgh, UK</i></p>
14:35 – 15:05 Networking/Exhibition/Refreshments	
15:05 – 16:50 Session Twelve: Immunology, disease and host-pathogen interactions	
<i>Moderators: Dan Macqueen</i>	
15:05 – 15:20	<p>TH-Oral-19 Co-diversification of salmonid hosts and its intestinal Mycoplasma</p> <p><i>Morten Limborg, University of Copenhagen, Denmark</i></p>
15:20 – 15:35	<p>TH-Oral-20 Haplotype-based association analyses to inform fine-mapping of QTLs linked to IHNV resistance</p> <p><i>Christopher Setzke, University of Washington, USA</i></p>
15:35 – 15:50	<p>TH-Oral-21 Cell specific differential expression of duplicated genes in Atlantic salmon liver</p> <p><i>Richard Taylor, University of Edinburgh, UK</i></p>



<p>15:50 – 16:05</p>	<p>TH-Oral-22 Gill health, microbial dysbiosis and husbandry interventions at sea <i>Benjamin Clokie, University of Stirling, UK</i></p>
<p>16:05 – 16:20</p>	<p>TH-Oral-23 Hologenomics of resistant/susceptible Atlantic salmon families to sea lice <i>Cristian Gallardo-Escárate, Interdisciplinary Center for Aquaculture Research, Chile</i></p>
<p>16:20 – 16:30</p>	<p>TH-Flash-17 Investigating the genetic basis underlying response against salmonid alphavirus in Atlantic salmon <i>Domniki Manousi, Norwegian University of Life Sciences, Norway</i></p>
<p>16:30 – 16:40</p>	<p>TH-Flash-18 MicroRNAs and their predicted target genes associated with winter-ulcer in Atlantic salmon <i>Rune Andreassen, Oslo Metropolitan University, Norway</i></p>
<p>16:40 – 16:50</p>	<p>TH-Flash-19 Supporting disease control in Atlantic salmon aquaculture using viral genomic surveillance <i>Albert Knight, The Roslin Institute and the Royal (Dick) School of Veterinary Studies, The University of Edinburgh, UK</i></p>
<p>19:00 – 23:00</p>	<p>Conference Dinner</p>