## Uncertainties in Modelling Water-borne Disease Transmission among Salmon Farms in the Discovery Islands, British Columbia

Mike Foreman<sup>1</sup>, Kyle Garver<sup>2</sup>, Dario Stucchi<sup>1</sup>, Ming Guo<sup>1</sup>, Darren Tuele<sup>1</sup>, Jared Isaac<sup>3</sup>, John Morrison<sup>1</sup>

<sup>1</sup>Institute of Ocean Sciences, Sidney

<sup>2</sup>Pacific Biological Station, Nanaimo

<sup>3</sup>Department of Mechanical Engineering, University of Victoria



Fisheries and Oceans Canada Pêches et Océans Canada



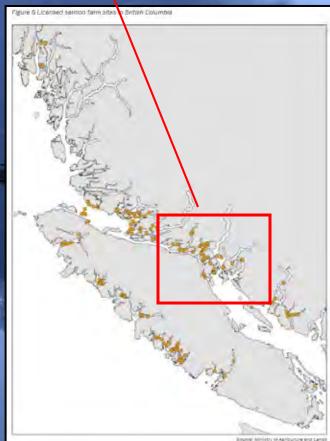




## **Outline**

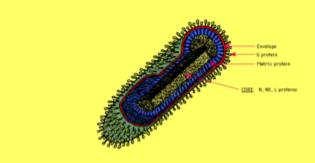
- Background & motivation
- Circulation model
- Biological model
- Uncertainties
- · Summary & future work





## Infectious Hematopoietic Necrosis Virus (IHNV)

- Infects a variety of salmon and trout species along the northeast Pacific
- BC farmed Atlantic salmon infections:
  - 1992-97: 14 netpen sites near Campbell River
  - •2001: 26 sites, spread from Campbell River
  - •2002: 10 sites on west coast of Vancouver Island
- High costs- culling entire farms





#### What was learned?

- Transmission spread rapidly
- · Infected farms had an identical virus type
  - → farm to farm transmission
    - Anthropogenic (poor bio-security)
    - Waterborne (needs more study)



# Model Components

#### 1. Circulation model:

- FVCOM
  - Finite Volume Coastal Ocean Model
- Requires observational data to initialize, force, & evaluate



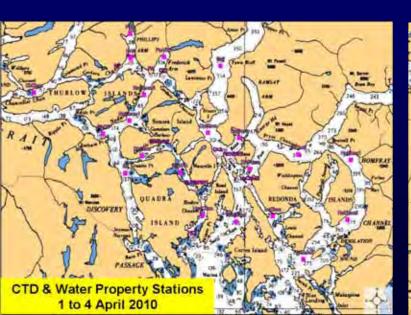
## 2. Coupled biological model:

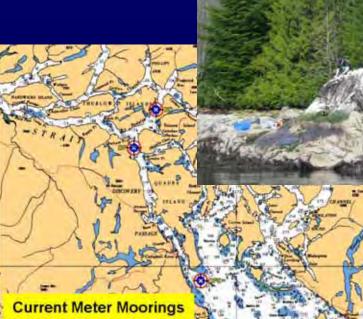
- Takes FVCOM output + UV radiation observations to disperse & kill I HN viruses originating on salmon farms
  - Standard "particle" tracking
- Requires lab experiment results to specify model parameters & relationships
  - Shedding rates, minimum infective dosages, virus stability, ...

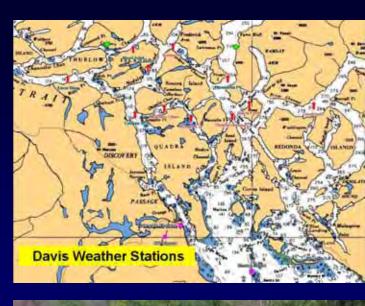


## Field Observations

- Weather stations:
  - 12 deployed: Oct 2009 to Feb 2010
  - Measure winds, UV radiation, heat flux components
- Water property surveys:
  - Temperature & salinity profiles
- Current Meter Moorings:
  - 3 deployed in Oct 2009

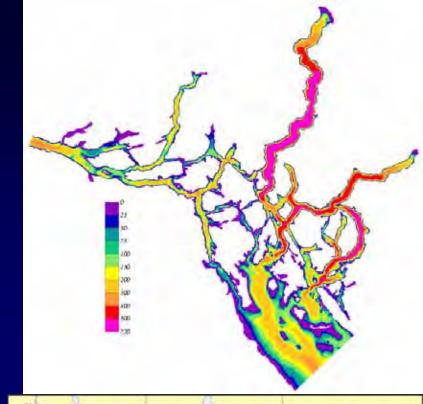






# Regional Challenges

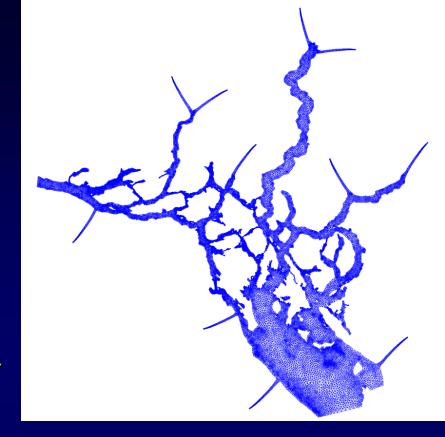
- Deep fiords with seasonal river discharges
  - Strong stratification
  - Potential baroclinic pressure gradient problems in FVCOM
- strong mixing in island channels
  - Some of the strongest tidal currents in the world

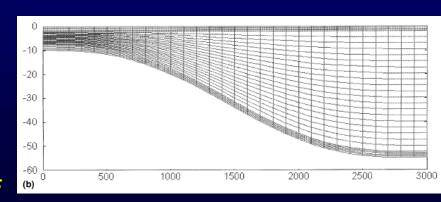




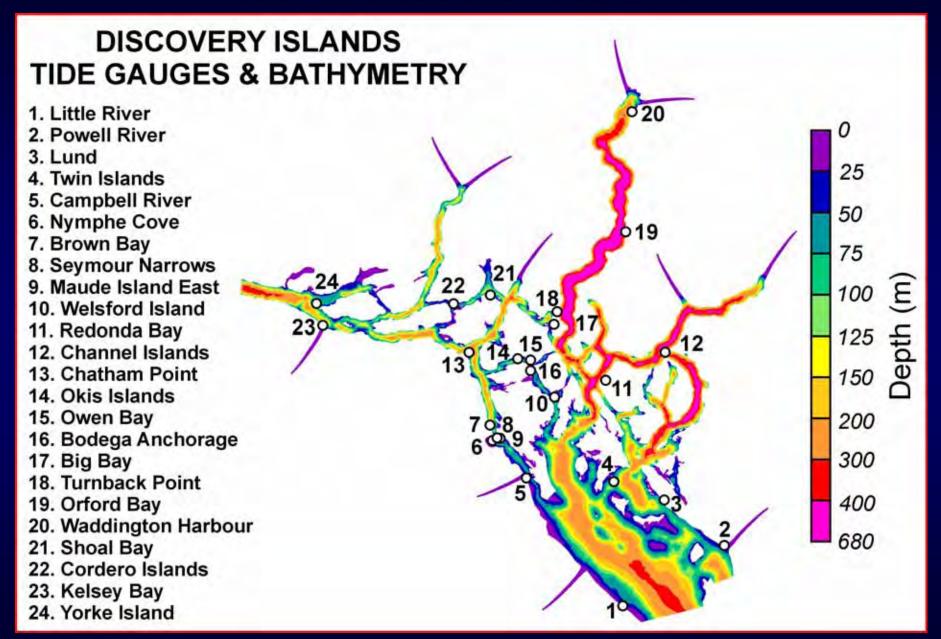
## Model Overview

- Simulation for April 1-28, 2010
  - Tides, river discharge & weather station data for forcing
  - April 1-4 CTD observations blended with climatology to initialize TS fields
  - Current meter & tide gauge observations to evaluate accuracy
- Horizontal grid:
  - 37596 nodes, 68467 triangles
  - Resolution from 1.7km to 90m
  - 11 rivers
- Vertical grid:
  - 20 unequally-spaced sigma coordinates

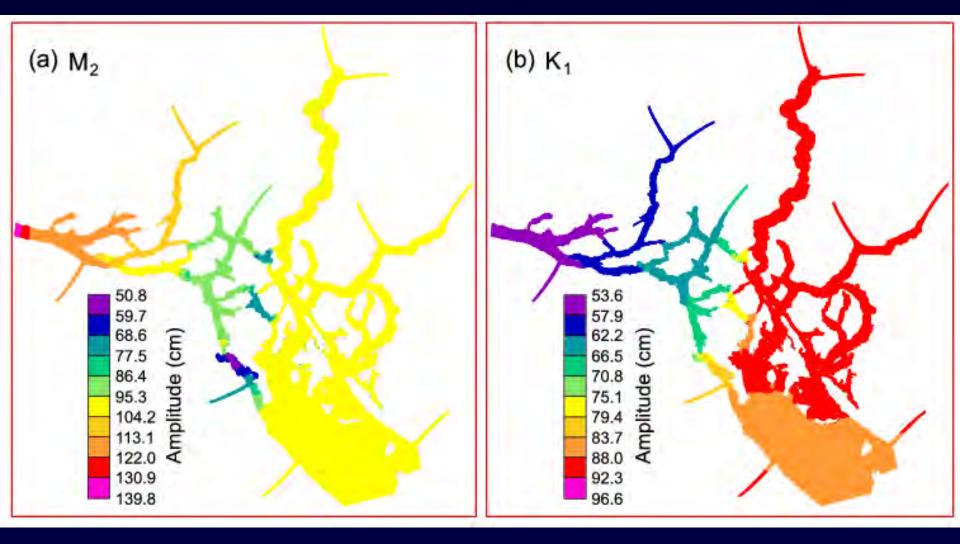




#### Model Evaluations: sea level



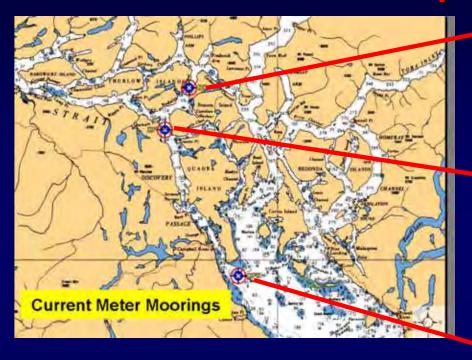
## M<sub>2</sub> & K<sub>1</sub> Tidal Elevation Evaluations



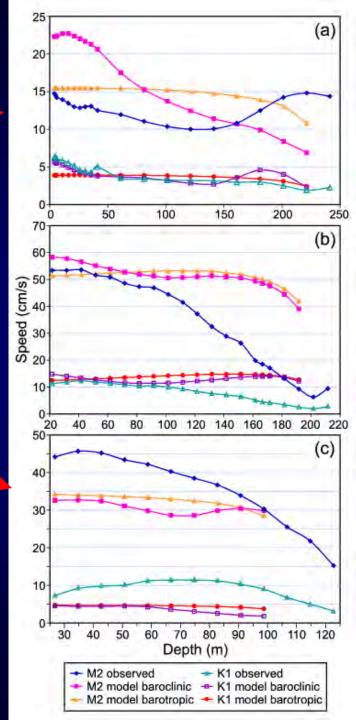
Average amplitude/phase errors vs 24 tide gauge locations

- distance in complex space
- M<sub>2</sub>: 3.9 cm
- K<sub>1</sub>: 3.2 cm

## M<sub>2</sub> & K<sub>1</sub> Tidal Current Evaluations vs Depth



- K<sub>1</sub> model speeds generally OK
- •M<sub>2</sub> not as good
  - Surface values good at Discovery but deteriorate with depth
  - off Cape Mudge, too weak
  - in Nodales, too strong

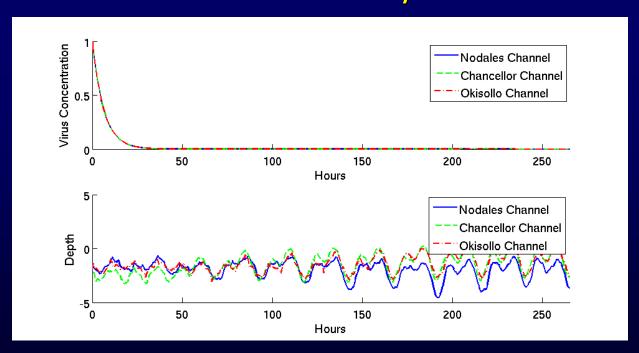


# IHN Virus Biological Model

IHN virus mortality function of T, S, UV radiation

$$V(t) = V(t-1) * \exp(-a * U(t)_0 * \exp(-kz) - b)$$

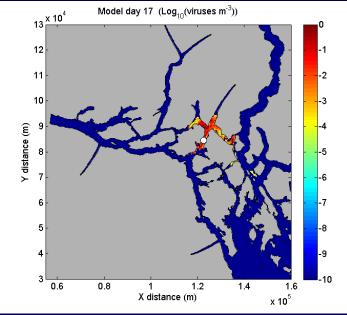
V(t) is the virus concentration, U(t) is the UV radiation value at sea surface, z is the depth, a, k and b are constant values determined from lab experiments

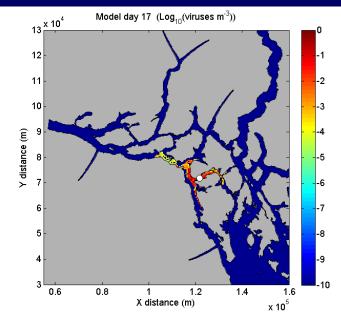


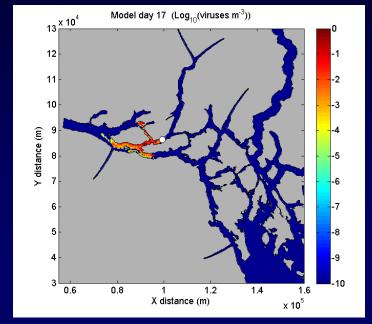
Time series of virus concentration and average depth for 3 farm releases

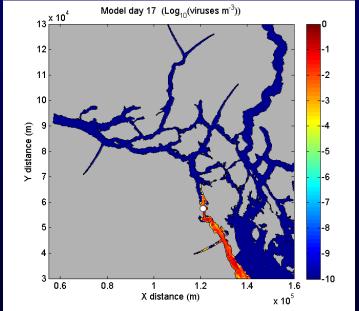
## Preliminary Near-Surface Concentrations

(100,000 virus particles released per hour over 19 days & in top 2m, tracked for 5 days)





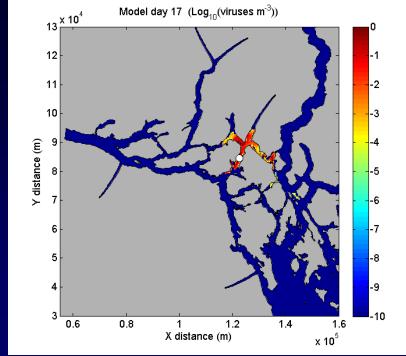




# Will nearby farms be infected?

- Model viral releases need to be scaled by the shedding rates
  - Lab work estimated 6 x10<sup>10</sup>
     viral particles per hour
  - Scale-up present concentrations (max 1 virus/m³) by 6x10<sup>5</sup>
- Model concentration fields need to be interpreted in terms of minimum infectious dose
  - Lab min dose is

    10 PFU/ml = 10<sup>7</sup> virus/m<sup>3</sup>
  - ➤ Nearby farms may not be infected (under present assumptions)





## Model Uncertainties

- Biological model
  - Extension of lab-based parameters & relationships to open ocean
    - Shedding rates; minimum infective dosages; virus stability with UV, T, S;
  - Neglect of other important factors (e.g., bacterial content)?
- Physical model
  - accuracy of model fields (u, v, w, T, S)
  - Sufficient range of model simulations (over different forcing conditions)





## Reducing Uncertainties

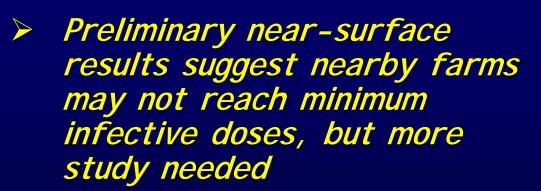


- Biological model
  - Field observations (where feasible) & more lab experiments
  - An actual disease outbreak to provide data
- Physical model
  - More observations to evaluate model accuracy
  - Improvements in model forcing fields, resolution, physical processes, ...



# Summary

Preliminary circulation & IHN virus water-borne transmission models have been developed for the Discovery I slands



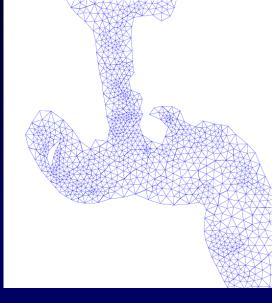
- Many uncertainties
  - Some can be reduced





# Future Work





- Model details:
  - Move to higher resolution grid
    - Better coastline & depths
  - Implement newer/faster(?) version of FVCOM
- Model simulations
  - Virus releases below 2m
  - include other mortality dependencies (e.g., T,S)
  - Later in summer with more freshwater

