Extending the Validity of a One-Dimensional Coupled Bio-physical Model by Parametrization

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Outline

- Strait of Georgia
- Our goals
- The model
- Example parameterizations
- Conclusions

Fraser River

Strait of Georgia

North-east Pacific Ocean

ohnstone

50 mi

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uan de Fuca

Strait of Georgia

Fraser River sockeye returns



Number of returning salmon (millions), 1952 to 2009

Note: it takes four years for most sockeye to return to spawn after hatching.

Biodiversity Canada who adapted it from LaPointe, 2010, with 2010 and 2011 added



Home waters for most of the population of BC.

Important habitat for juvenile salmon.

Important aquaculture centre.



Qualicum Bay Scallops

Strait of Georgia

- 400 m deep, 120 km long, 30 km wide
- significant input of freshwater
- strongly stratified, often without a surface mixed layer
- estuarine type circulation
- biologically productive



Johnstone

Northern Strait of Georgia

Traffic

Southern Strait of Georgia

Juan de Fuca



Satellite

Fraser River

Map

Hybrid



after LeBlond 1983 with Masson 2004 and Pawlowicz 2007



Phytoplankton



One celled organisms that form the base of the food chain in the ocean Plankton: can't swim against the currents

www.seriestemporales-ieo.net www.dnr.state.md.us www.nies.go.jp





Phytoplankton



Need light and nutrients to grow. Limiting nutrient: Nitrogen

www.seriestemporales-ieo.net www.dnr.state.md.us www.nies.go.jp









Halverson & Pawlowicz, 2012

Goal

 An accurate physical model to provide a foundation for lower trophic layer models (questions of primary productivity, timing and interannual variability) and for a carbon model (questions of pH values and processes)



Spring Bloom along the Ferry Route



Mark Halverson



One-dimensional Mixing Model plus

- Using a 1-dimensional model allows detailed vertical resolution and runs quickly; allows multiple parameter tests, grad-student friendly
- All two-dimensional processes must be parameterized
- Use a standard, well supported mixing layer model (KPP)
- Add baroclinic pressure gradients (due to finite size of the Strait)
- Add estuarine circulation





Vertically exaggerated. 40 m deep separated into 0.5 m disks (all the way down)





Physical Model

- Initialized with a temperature and salinity profile in Sep/Oct
- Forced with hourly wind, temperature, humidity, cloud fraction and daily Fraser River and Englishman River flows
- Bottom boundary condition: seasonal cycle of salinity and temperature and observed temperature increase over time



Estuarine Circulation



FIG. 2.7. Estuarine circulation in a typical British Columbia i let. Salt water entrained and carried seaward by river outflow is replenished by a net inflow at depth. Sloping isobalines (lines of equal salinity) indicate a down-inlet increase in salinity in surface brackish layer. Turbulent mixing occurs in vicinity of sill.

Entrainment of deeper, nutrient-rich, water into the surface outflowing circulation is the most important source of nutrients to the system.

Physical Model Major unresolved process: Estuarine Circulation



Amount of vertical entrainment set by conservation of mass and salt using data.

Parameterization Philosophy

- Only uses information available, either from the incoming forcing data (e.g. river flow) or from the core model (e.g. mixing layer depth)
- Is physically based
- Matches the observations
- Is numerically appropriate (i.e. No pathological behaviour at extrema)

Needed:

Mechanism:

- Amount of freshwater
 Salinity fit to add
- Depth to add
 Freshwater fit freshwater
- Vertical velocity rate
- Entrainment calculation

Parameterize Based on Data #1 Amount of Freshwater

- 48 cruises to the SoG
- Surface salinity is related to total freshwater input to the SoG (fit 1)
- Amount of freshwater is tuned to give the fitted surface salinity



Fit 1



Parameterize Based on Data #2 Depth of Freshwater

- Model calculates the mixing layer depth
- Freshwater is introduced (exponential) over a depth proportional to the mixing layer depth
- Proportionality is tuned to match the halocline depth between model and observations



A Salinity Similarity Solution

- Freshness ($F = S_D S$) almost has a similarity solution with depth S_D is deep salinity, S is in-situ salinity
- Freshness varies with distance from the river mouth and with the amount of freshwater
- We assume these dependencies are partially separable
- Integrated freshness

$$I_{F} = \int F dz = F_{*} \mathcal{F}\left(\frac{z}{d}\right) \mathcal{F}_{r}\left(\frac{r}{r_{3}}\right) \mathcal{F}_{T}\left(\frac{T}{\overline{T}}\right)$$

where d(r, T), r_3 and \overline{T} are relevant scales for the depth, lateral distance and freshwater input, respectively.





0

0.5

1

P 1.5

2

2.5

3



Vertical Velocity

- Can use Knudsen's relations (conservation of mass and salinity) to relate *I_F* to the entrainment velocity at depth
- By matching freshwater variation, one can show that the vertical velocity must vary with

$$\left(\frac{T}{\overline{T}}\right)\mathcal{F}_{\mathcal{T}}^{-1}\mathcal{D}_{\mathcal{T}}$$

where $\mathcal{D}_{\mathcal{T}}$ is the variation of the depth d with freshwater flow.

• I have no measurements of vertical velocity, but do have salinity profiles and can thus determine the freshness solution and can calculate this function versus total freshwater, *T*.

Entrainment Velocity vs T



Comparison to Riche & Pawlowicz

- Observation analysis shows entrainment almost independent of river flow
- This is consistent with the fit
- A constant value is wrong at large and small river flow, fit is forced to zero

Parameterize Based on Data #3 Entrainment Magnitude

- Using
 - the variation of entrainment with depth
 - the salinity fit (fit 1)
 - assumptions on the spreading shape of the plume = cone (1/3 to 1/4 of full circle) plus extra r⁰² (Hetland and MacDonald, 2008)
- One gets a magnitude of entrainment from 0.5-1.5 x 10⁴ m s⁻¹
- Agrees with box model of Riche & Pawlowicz

Parameterize Based on Data #3 Entrainment Depth Variation

- Assume a balance between the pressure gradient force and mixing of momentum
- Assume parabolic form _ℵ for 1/eddy viscosity
- Gives a cubic form for the entrainment.





Ferry chl fluor., Surface chl > 20 m, Surface model chl



Ferry chl unscaled between years

May 15, 2007

Results: Freshwater and Entrainment

- Adding the freshwater flow and entrainment allowed the simulation of the physical structure of the flow
- Biological code has been successfully used to hindcast and forecast the spring bloom timing in the Strait of Georgia.

2012 Spring Bloom



Conclusions

- Most models cannot resolve all important details
- We can "correct" these models by inserting our knowledge of processes but this requires:
 - Good knowledge of the processes
 - Careful thought for extremes
 - Lots of data to accurately tune parameterizations



Too much summer productivity



Another Water Mass



- Summer water mass advected from northern Strait has been near surface and so warm and low in nitrate
- Process probably happens throughout the year but has the greatest impact in the summer



- Amount of horizontal advection proportional to upwelling by Knudsen's relations
- Five f tting parameters: strength, time from surface, depth, width upward, width downward

Tuning Northern Advection

- Use temperature data from STRATOGEM to evaluate a given set of parameters (root mean square distance)
- Choose ranges for each parameter based on science
- Randomly choose 20-sets of parameters within the range
- Evaluate, choose top 6, combine & mutate, repeat



- Strength = 0.83 x Knudsen's scaling estimate
- Time from surface = 4 day back average
- Depth = 26 m
- Width Upward = 13 m
- Width Downward = 8 m







Results: Northern Return Flow

- Summer productivity and probably carbon cycle estimates require accounting for water mass entering at 20-30 m depth
- This water mass probably comes from the north and carries water that has been near the surface and is thus warm and low in nutrients
- Including this water mass, and tuning, reduces the temperature, nitrate, oxygen and carbon error between the model and the observations.

Conclusions

- Most models cannot resolve all important details
- We can "correct" these models by inserting our knowledge of processes but this requires:
 - Good knowledge of the processes
 - Lots of data to accurately tune parameterizations
- With biology included in the model, there are more cross-checks possible which both
 - Expose missing processes
 - Confirm parameterizations