



# Retentive structures, transport and connectivity in coastal ecosystems: using a quantitative particle tracking metric to describe spatio-temporal patterns

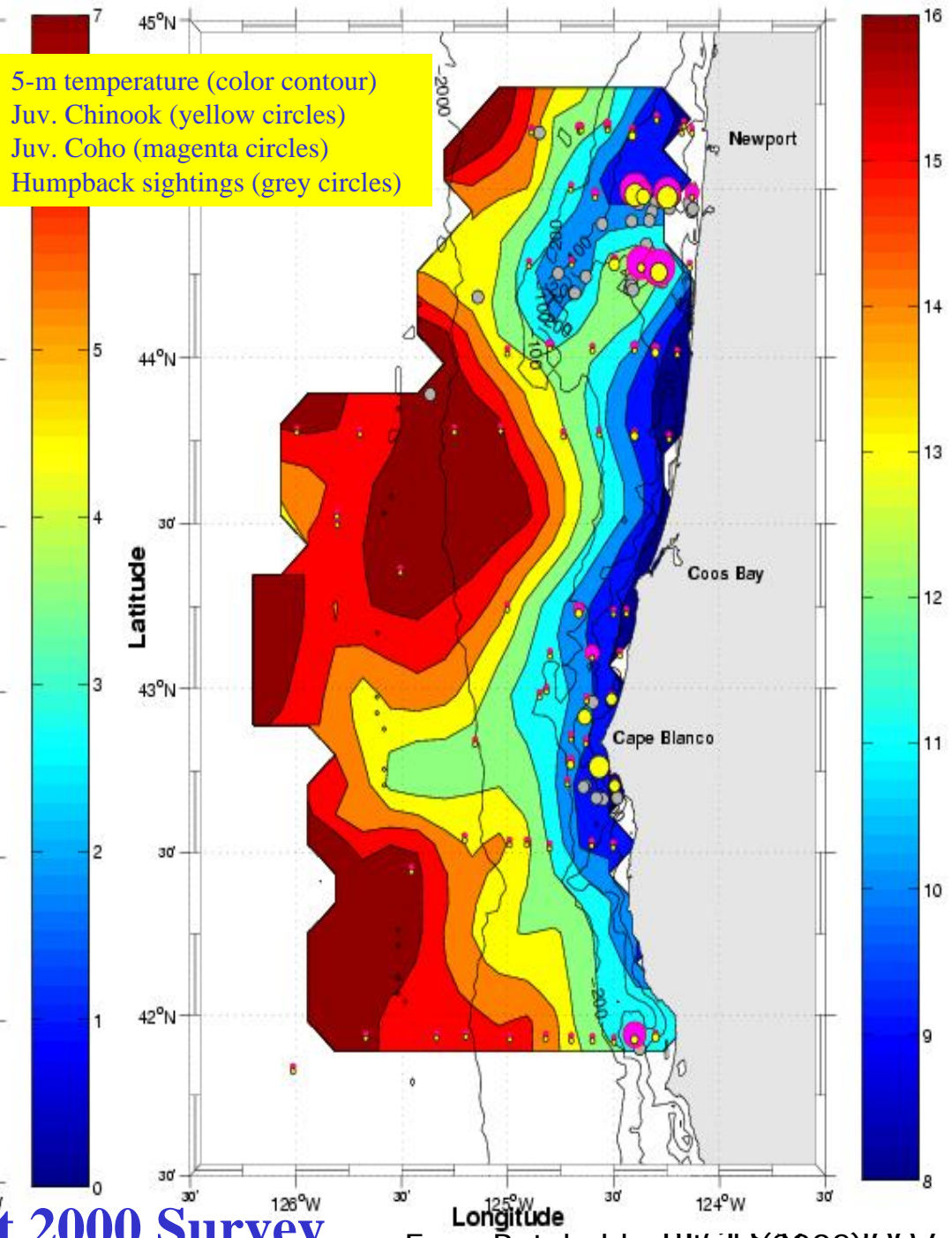
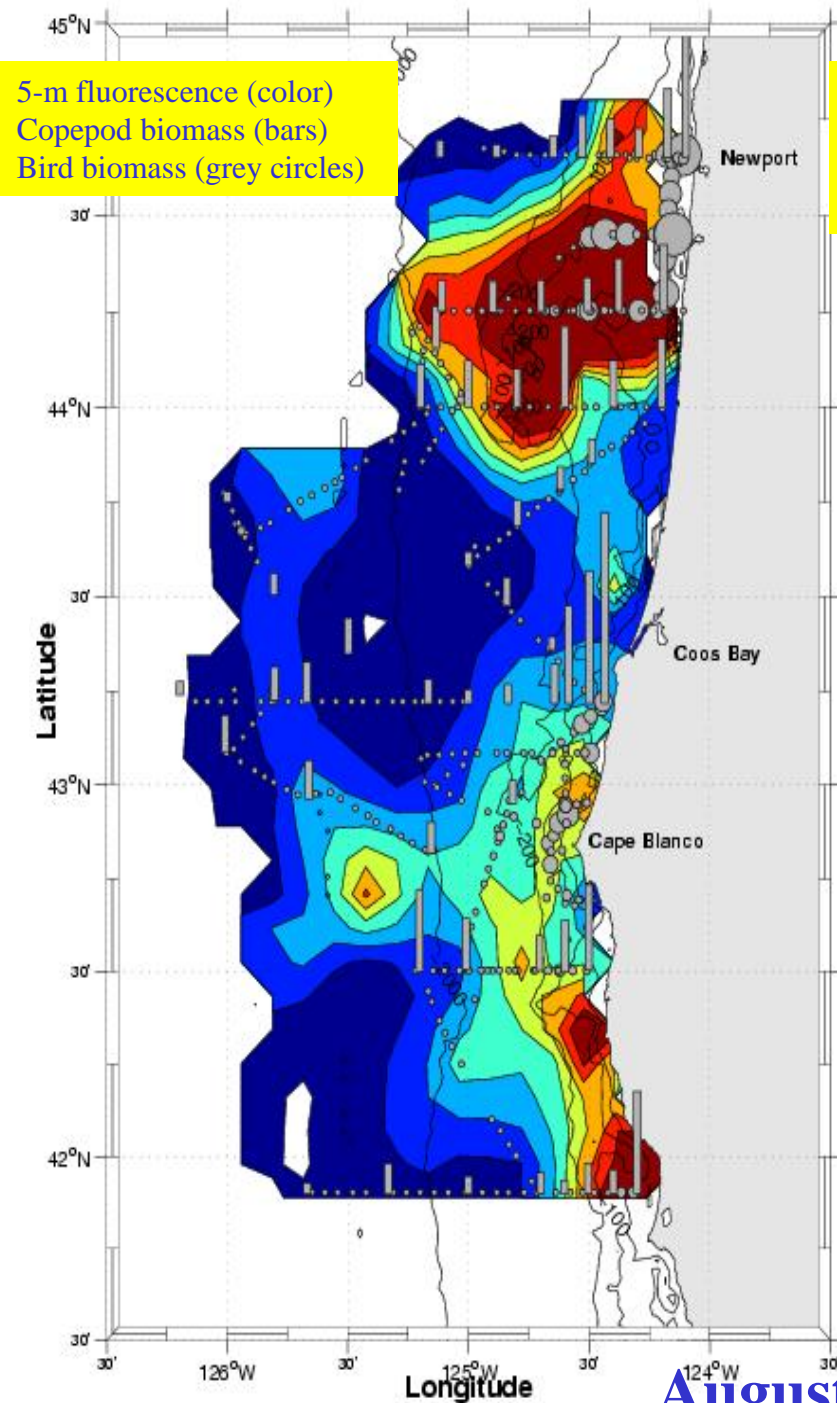
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## Roadmap

An introduction to what we are doing and why.

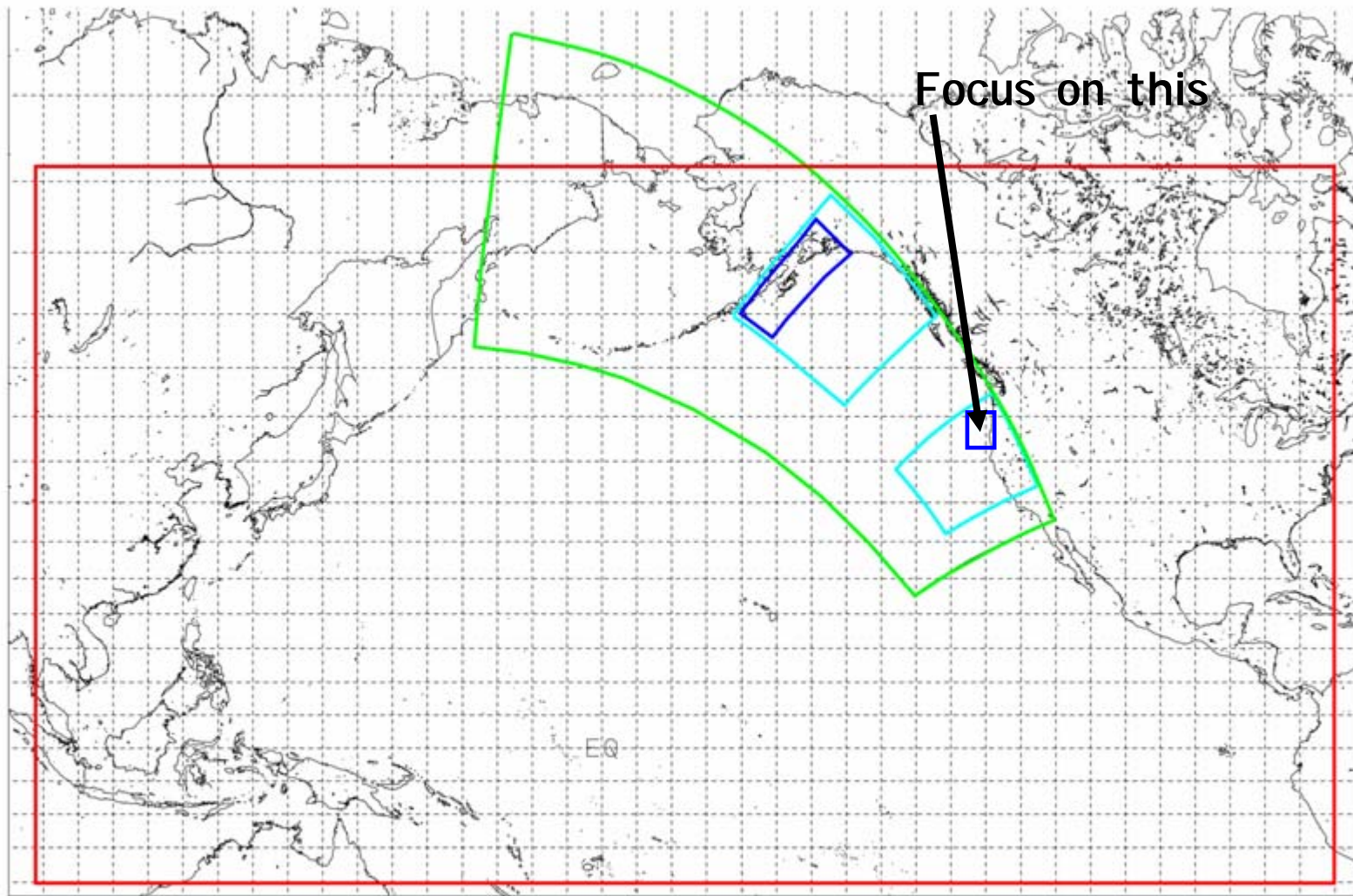
Methods

Conclusions/Summary



# Mesoscale Structures

- 10's to 100's of kilometers
- Persistence of weeks to months
- Have strong energetic interfaces; physical energy augments trophic energy of biological system (Bakun)
  - Biological production
  - Life cycle closure (larval retention sensu Sinclair)
- Associated biological structures on similar spatial and temporal scales
- Types: upwelling fronts, river plume fronts, shelf break fronts, eddies



Delta x = 20 40 km

Delta x = 10 km

Delta x = 3 km

Delta x = 1 km



# Implementation

## RCCS

Domain: 41 – 45.5N, -126.7 – 123.5E

ROMS: 166 x 258 x 42 gridpoints  
(~ 1 km res.)

COAMPS wind forcing; Blended product using 9-27 km resolution, but mostly 9 km over RCCS domain

Initial/boundary conditions provided by NEP model simulation from 2002.

Forward run for 2002

Daily averaged physical snapshots of velocity, temperature, etc.

## NEP

Domain: 20 - 73N, 115 – 210E

ROMS: 226 x 642 x 42 gridpoints (~10 km res.)

Subdaily (6 hr) T42 CORE wind and fluxes (Large and Yeager)

Initial/boundary conditions provided by CCSM-POP hindcast model

Forward run for 1958-2004— includes multiple El Nino's, Regime Shifts, and 2002 cold intrusion

Daily averaged physical snapshots of velocity, temperature, etc.

# Typical horizontal velocity field

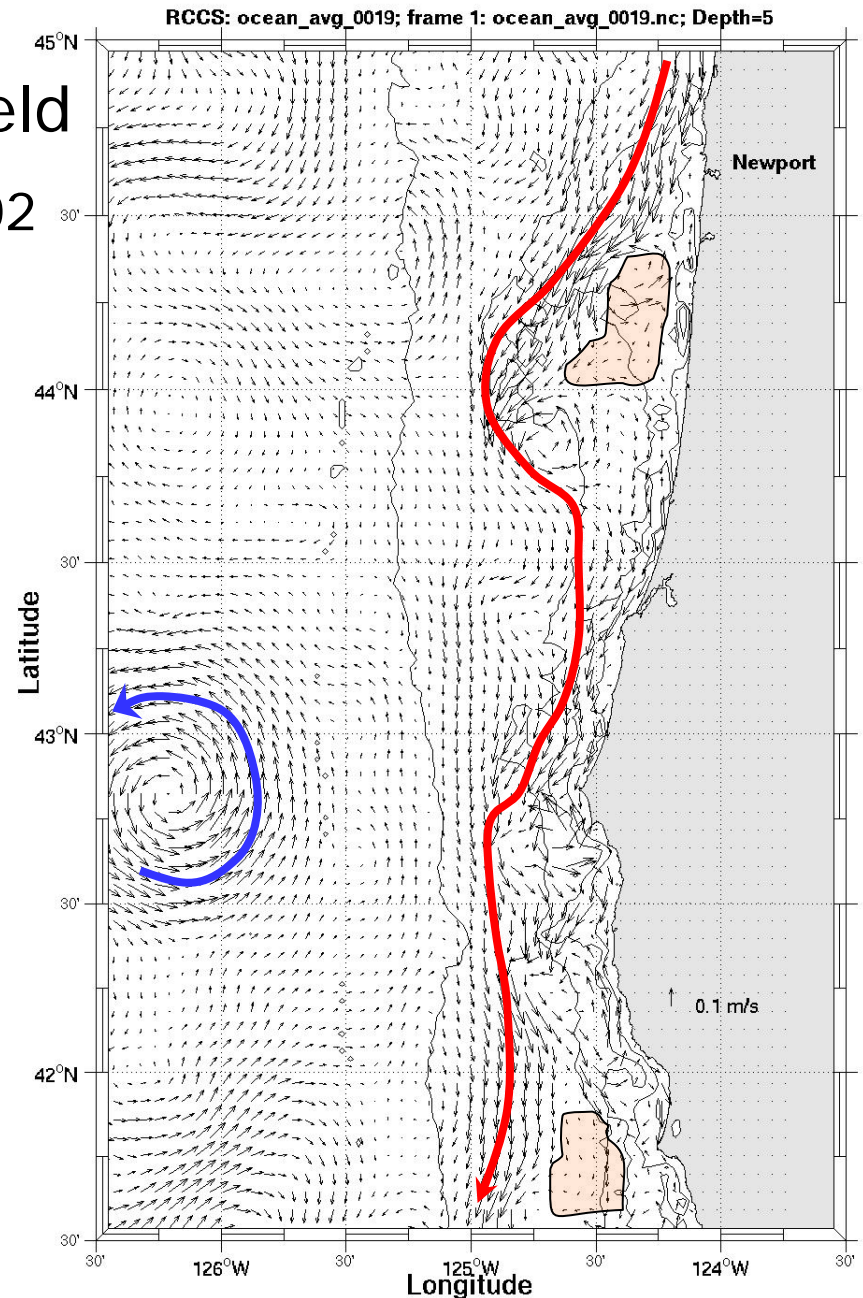
This one is from 5m on 1 July 2002

Alongshore upwelling jet

Large offshore eddy

Sluggish flow

Note: only every 3<sup>rd</sup> vector is plotted in both LON and LAT directions.



50,000 initial locations randomly selected

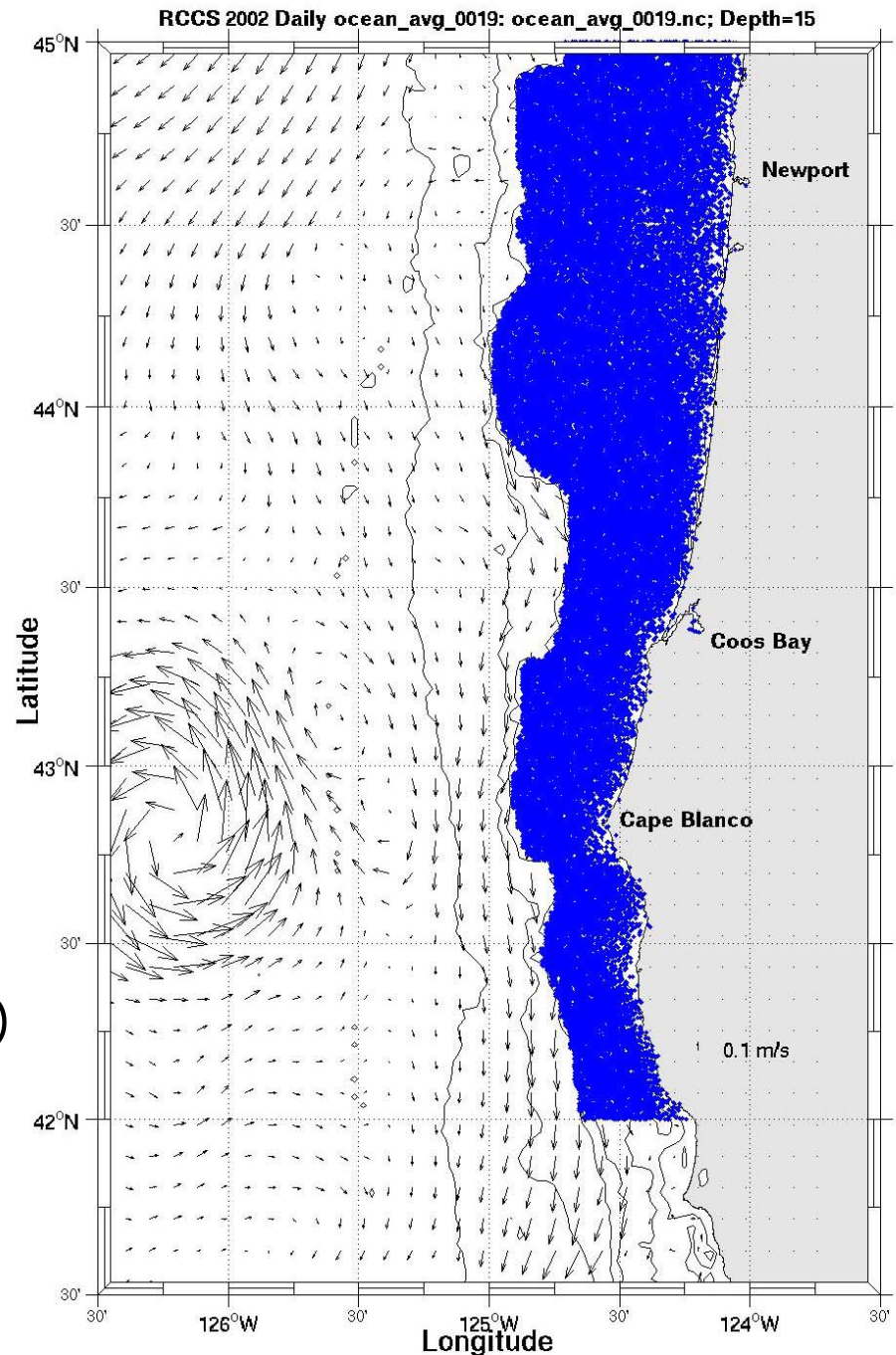
$70 \leq \text{Bottom depth} \leq 500$  meters

Latitude = 42-45N

(Averages ~ 1-2 indiv/km<sup>2</sup>)

Depth of particle was randomly assigned to be within 10-100m

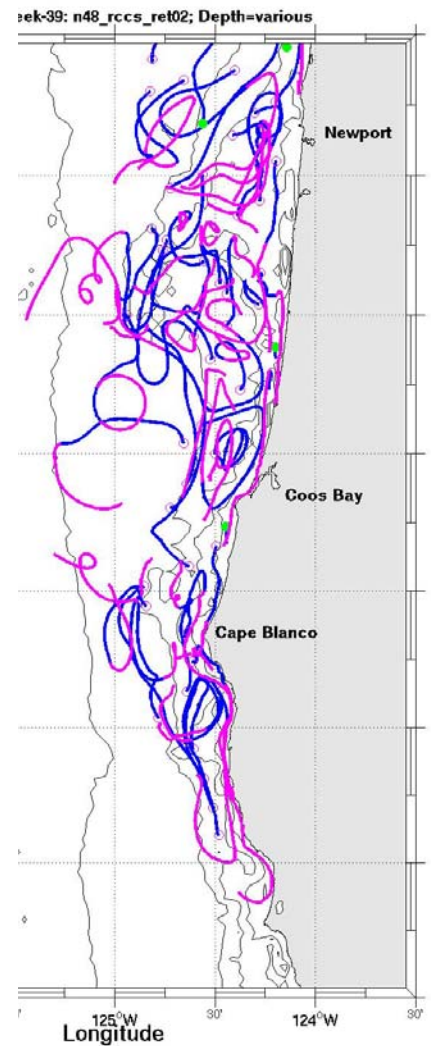
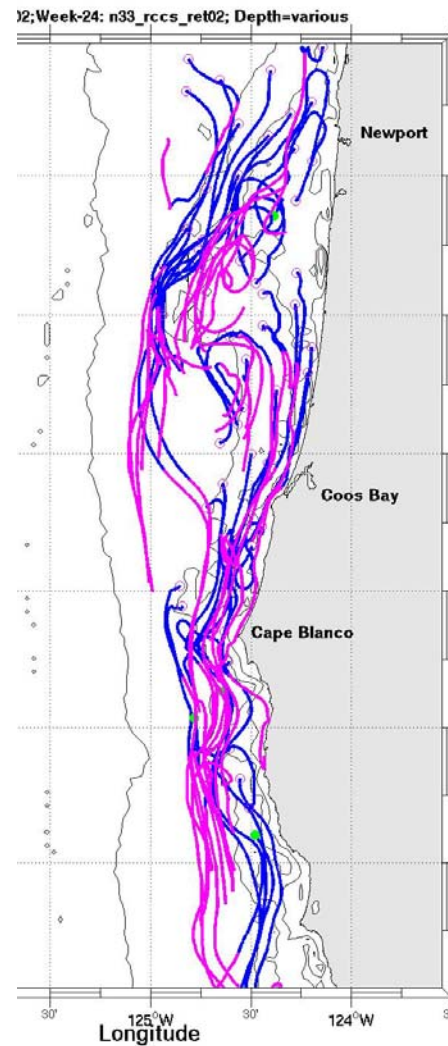
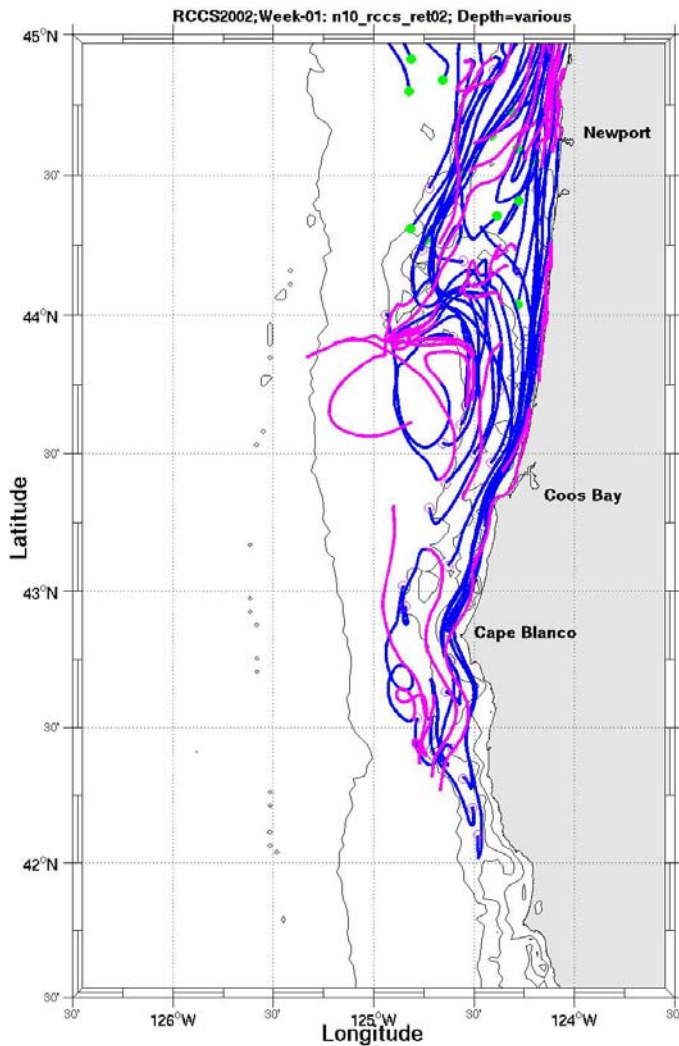
Beginning on 2 Jan 2002, each particle was 3D-advected for 15 days based on trilinear spatial & linear temporal interpolation of velocity fields with  $dt=1$  hr; a new simulation (same starting locations) was begun every 7 days for all of 2002 (total of 53 simulations)



2 Jan

12 Jun

25 Sep



Subset of 50 of 50,000 particles selected for plotting

Circle (start loc); Blue (ET=0-7.5d); Magenta (ET=7.5-15d); Grn circle (grounded)



# Untangling spaghetti . . .

## Retention Indices and Metrics

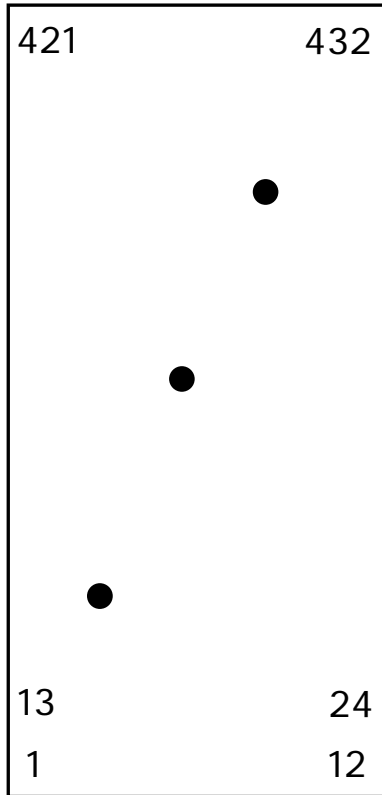
- Displacement distance at some elapsed time
- e-flushing time for a specified control volume (distance)
- retentive vortices (size, location, persistence)

## Connectivity Indices and Metrics

- Transition Probability Matrix Plots
- Sources and Destinations

# Retention Indices and Metrics

## Calculation Basics



Indices and Metrics require **gridding** of continuously distributed data.

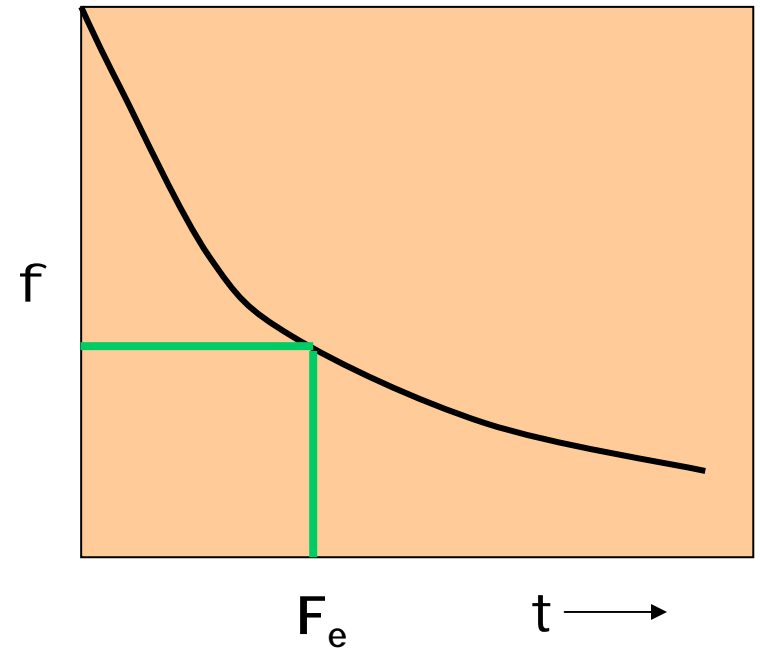
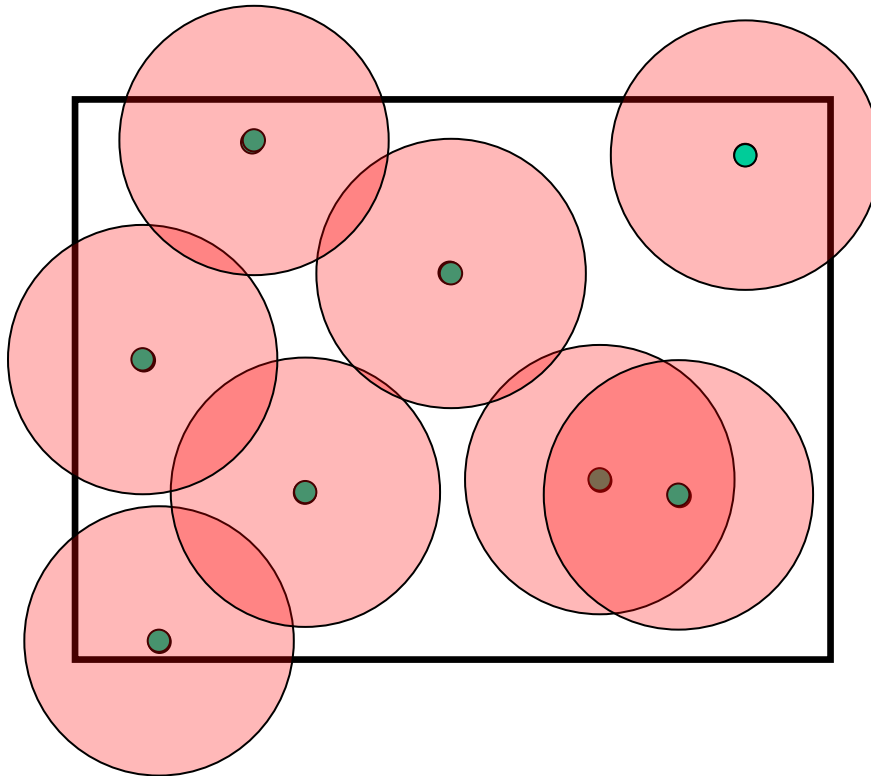
For this domain, the  $3^\circ$  of Latitude and  $1^\circ$  of Longitude were gridded into 5' bins, producing a display grid of 12 x 36. (432 cells total)

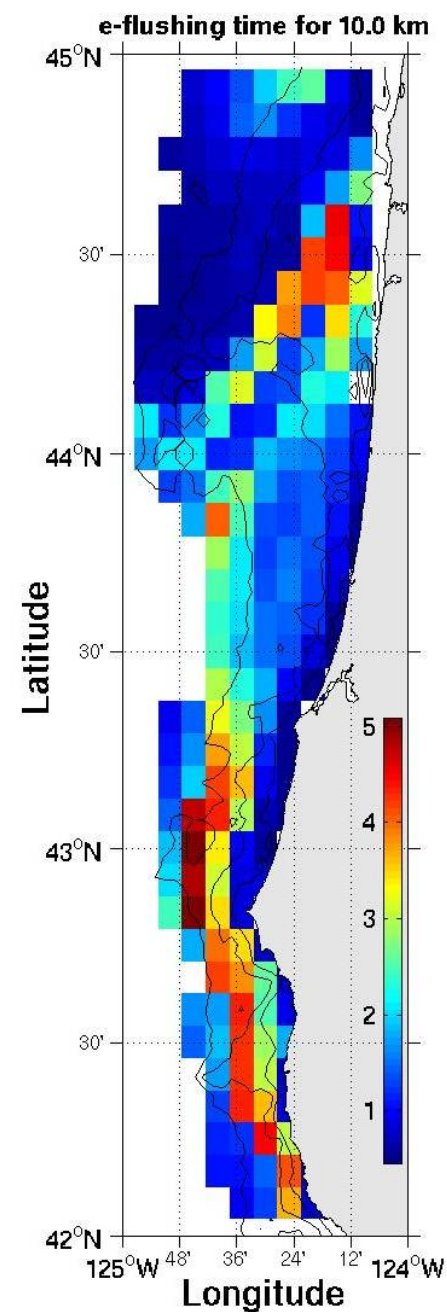
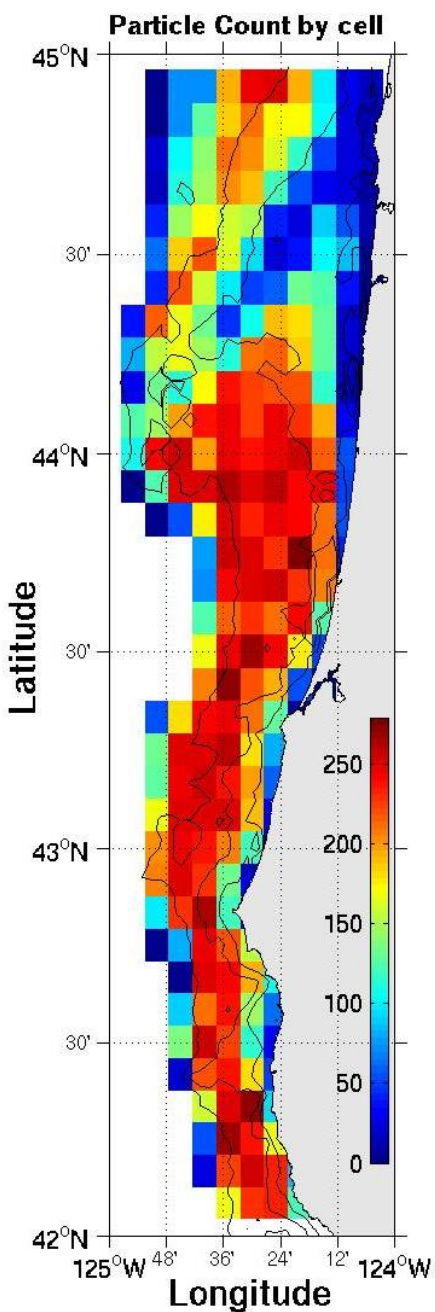
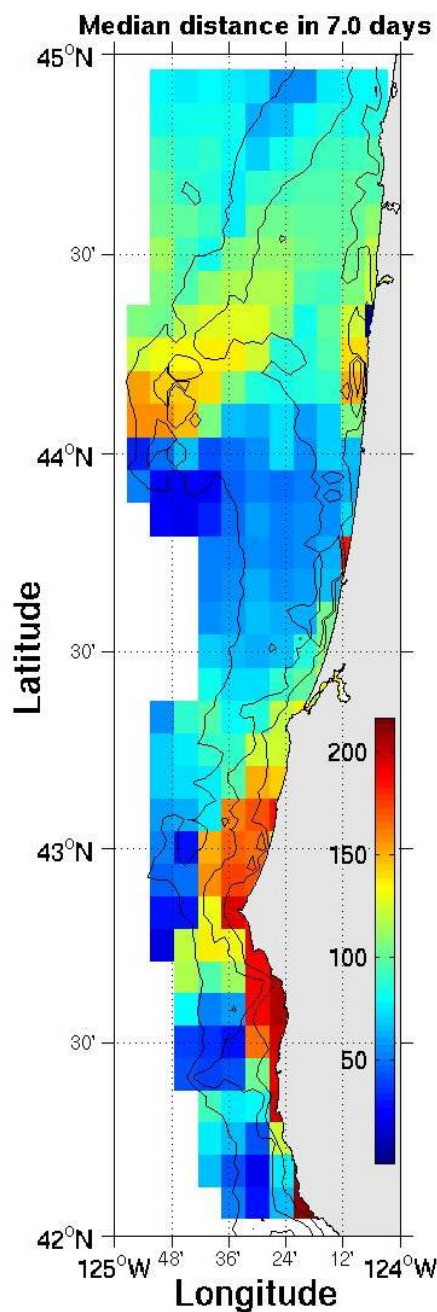
**Displacement:** Straight-line distance from origin to location at some later fixed  $\Delta t$ ; median, mean, std; usually eliminate particles with boundary interactions from statistics ( $\Delta t$  dependent)

**E-flushing time:** Time for a fixed fraction of particles within a control volume (or distance) to be lost from the volume ( $\Delta d$ ,  $\Delta V$  dependent)

# E-flushing Time ( $F_e$ )

- select a distance ( $r$ ) or control volume
- track fraction ( $f$ ) remaining within  $r$  of initial location as function of time ( $t$ )
- note time when  $f$  declines to  $< 1/e$  ( $\sim 0.368$ )

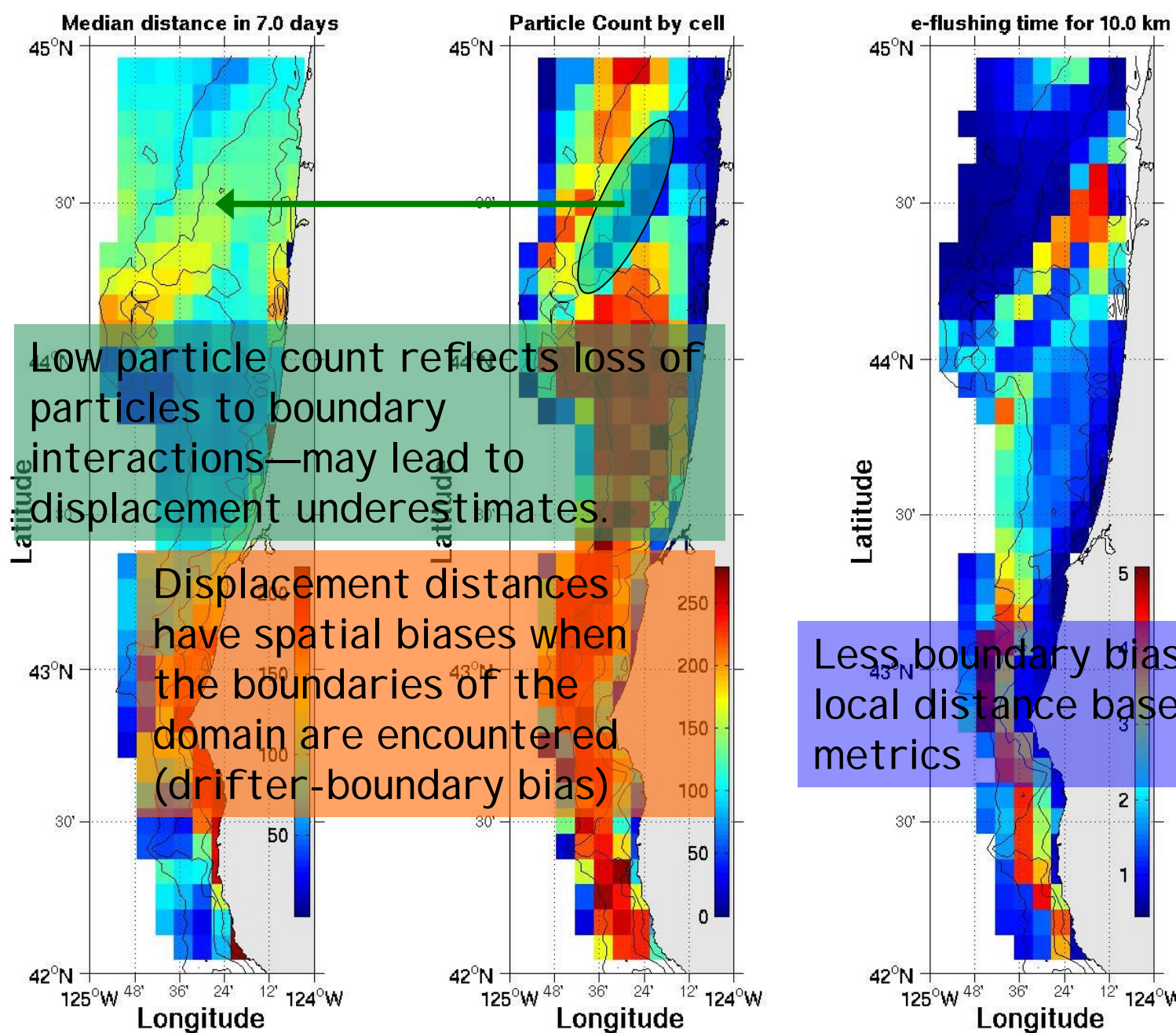




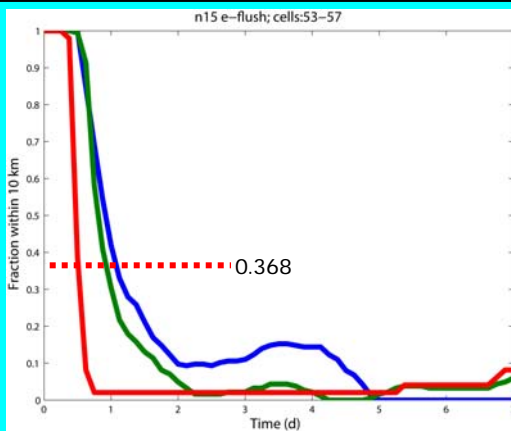
RCCS  
2 Jan

Start: 2 Jan 2002; Period of N flow; Particle counts are higher for initloc in south and lower velocity regions

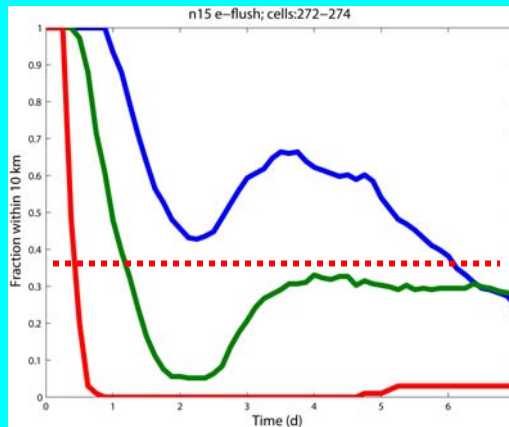




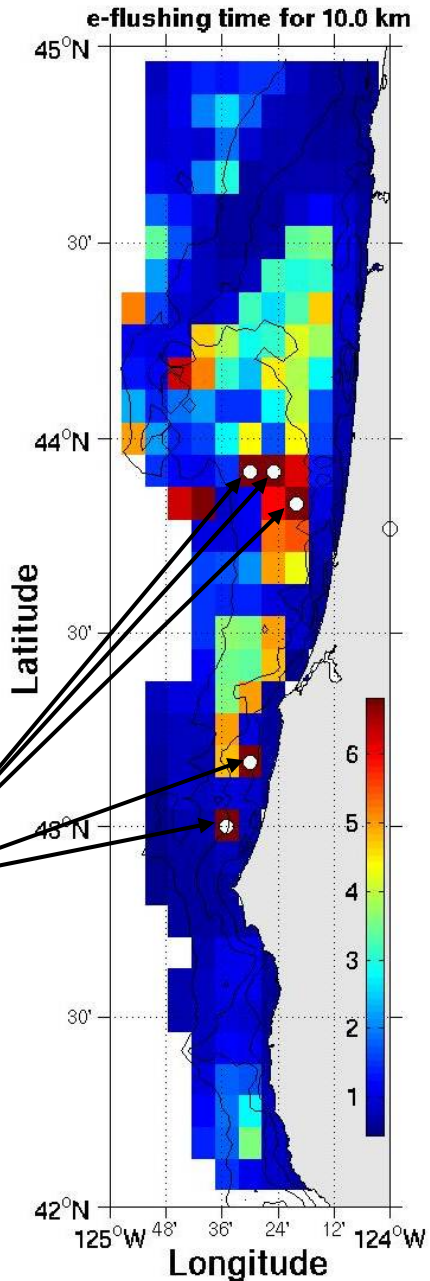
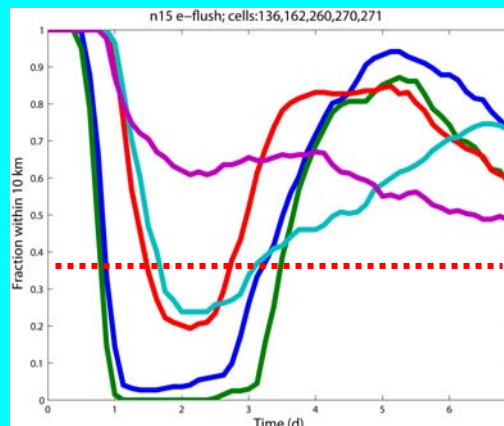
RCCS  
2 Jan



Well behaved  
"flushing" with  
single transition  
through  
threshold



Number of threshold  
crossings not ONE;  
rather 0 or >1; zero =  
slow, but regular  
flushing; >1 indicates  
substantial  
recirculation

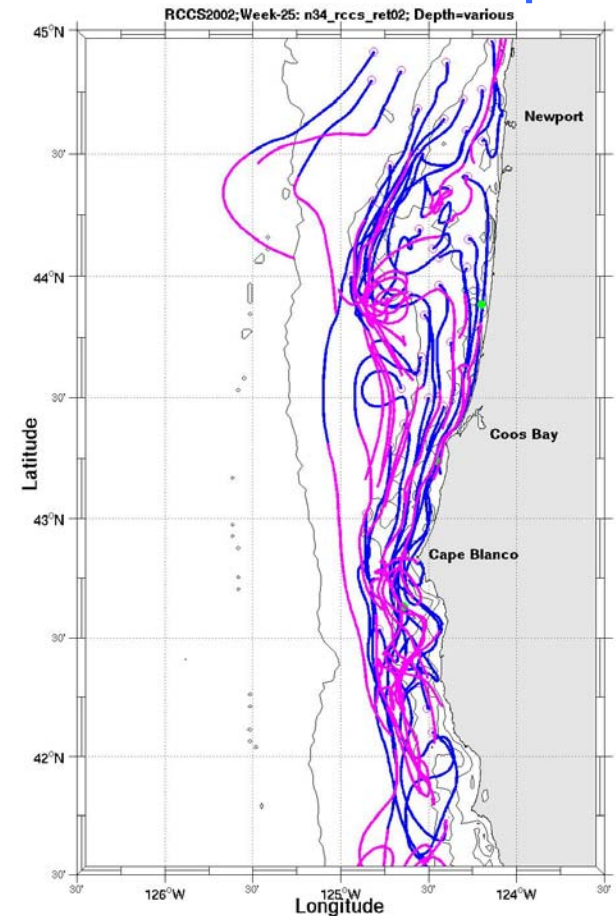
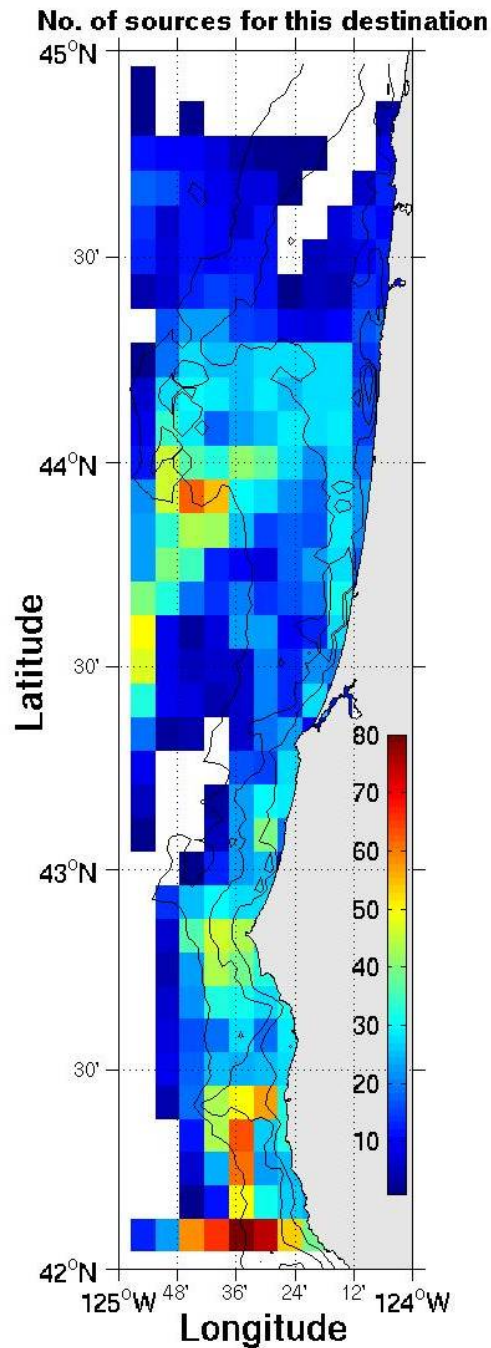
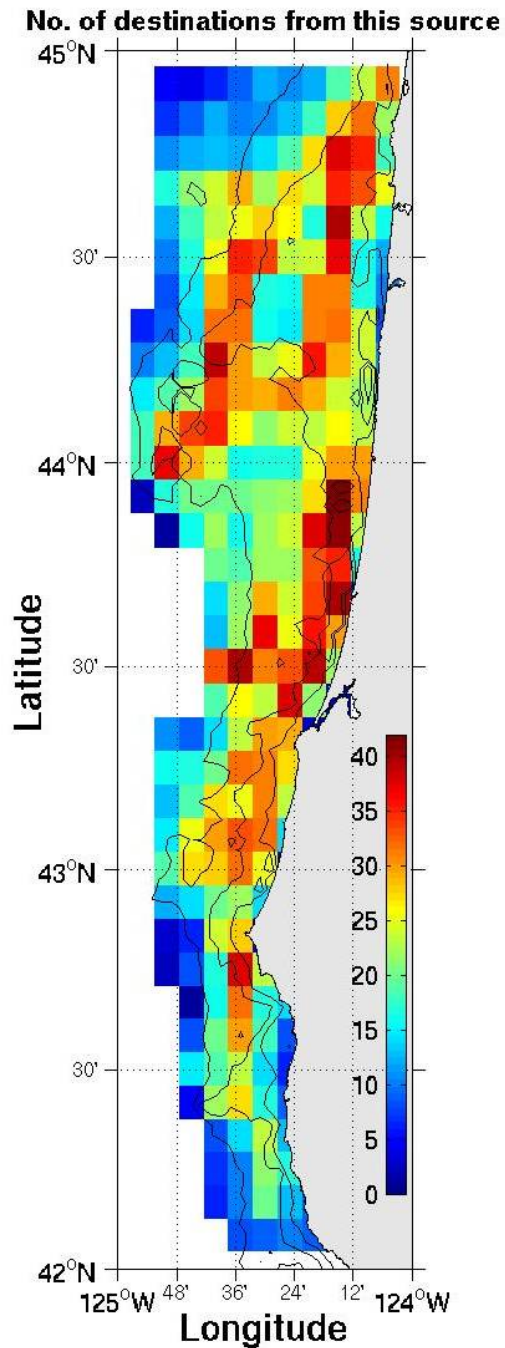


RCCS  
6 Feb

RCCS  
19 Jun 2002 start

ET = 7 days

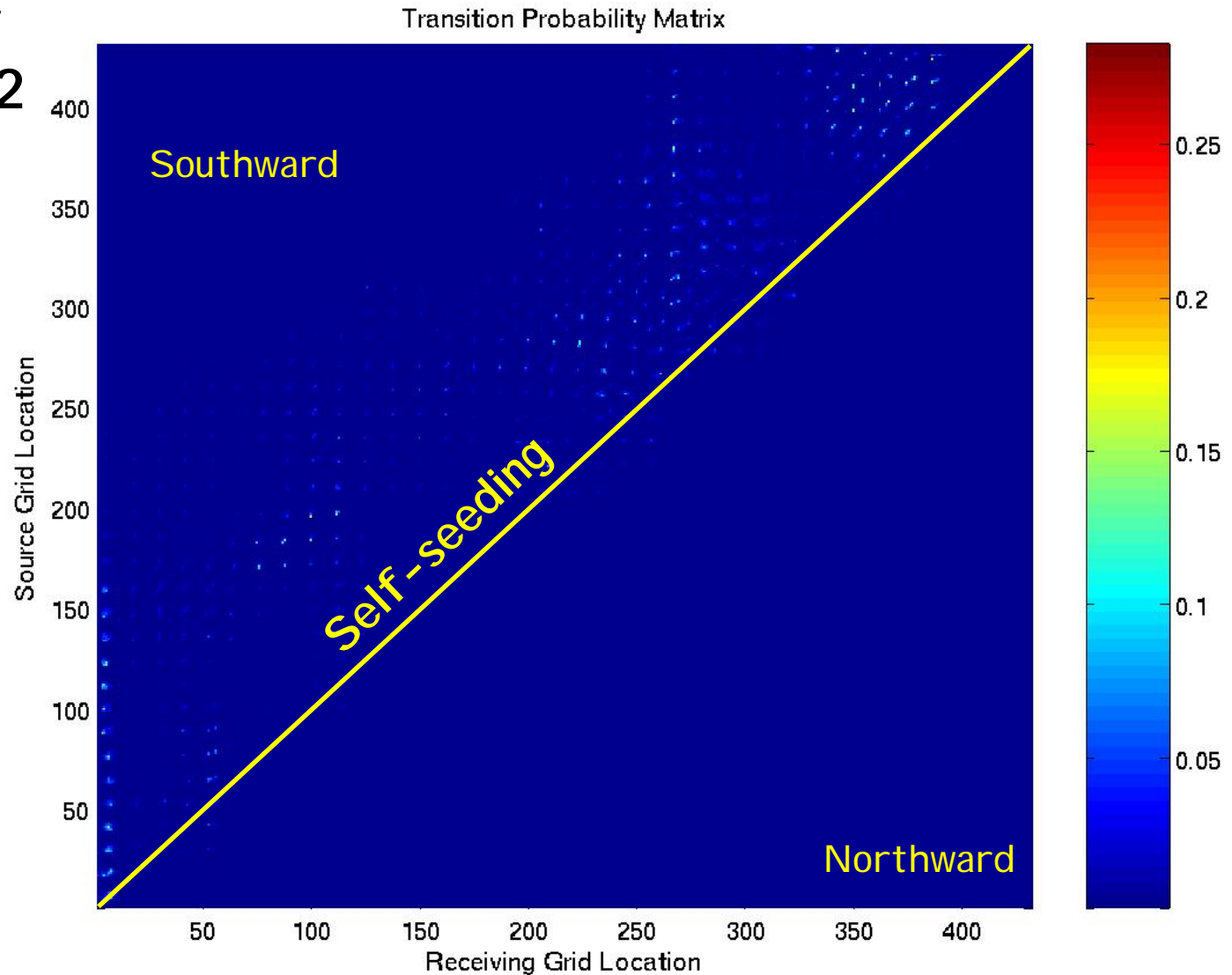
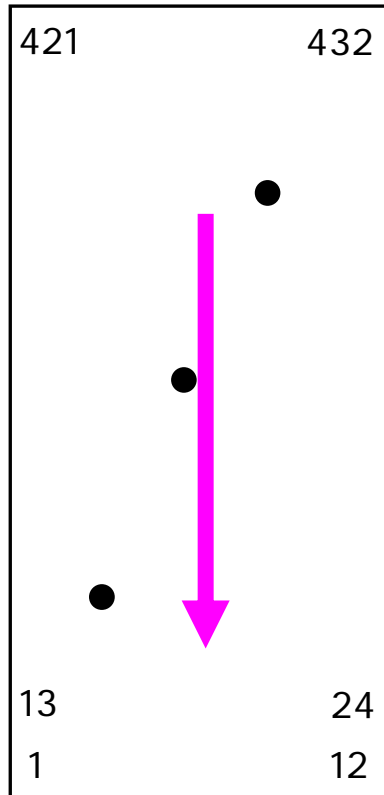
Strong Upwelling;  
southward transport





# Connectivity

19 Jun 2002

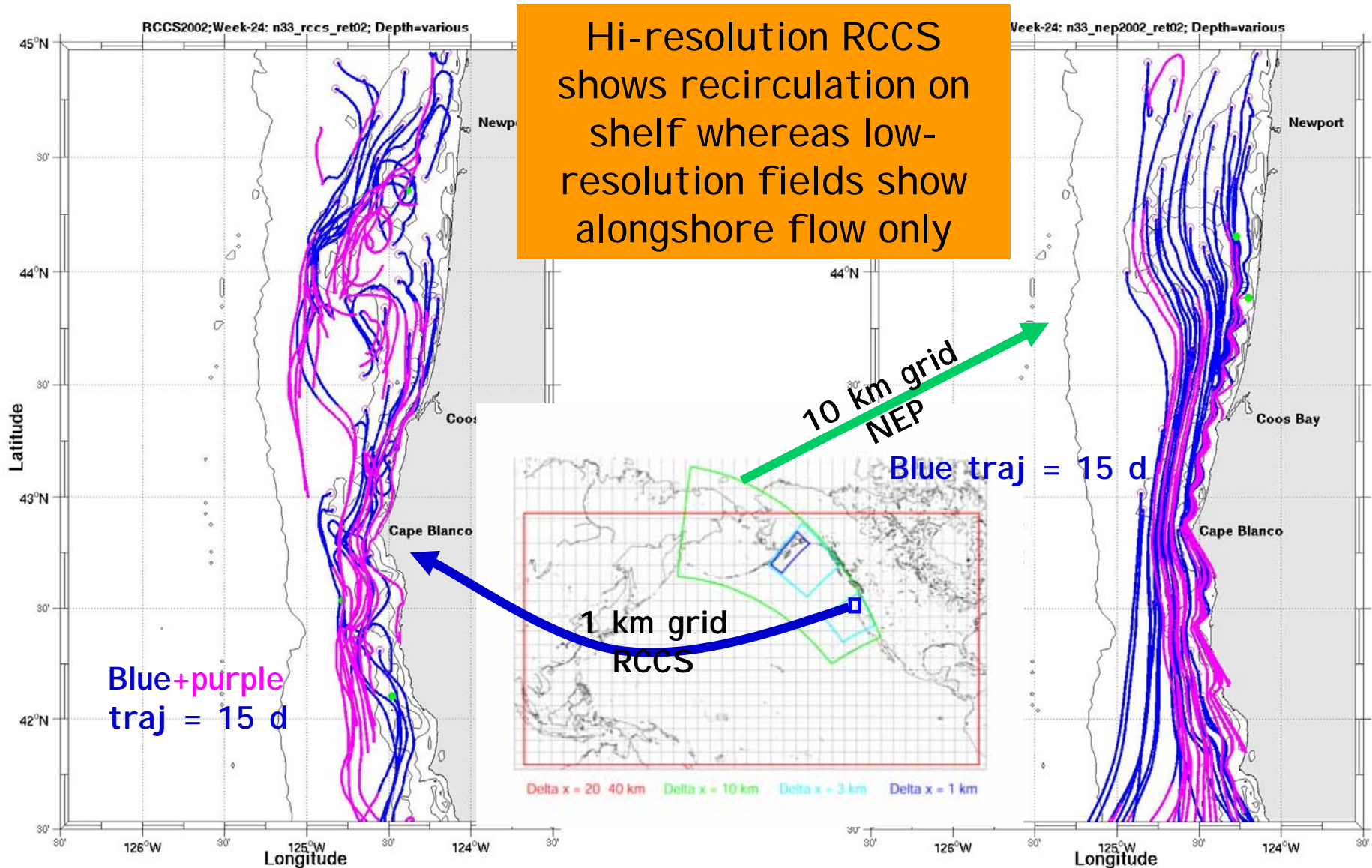


Probability that an indiv starting in grid cell (Y-axis) will be located in grid cell (X-axis) after 7 days. Numbering of grid cells is by row from lower left (SW) of domain to upper-right (NW). This shows southward movement of particles.



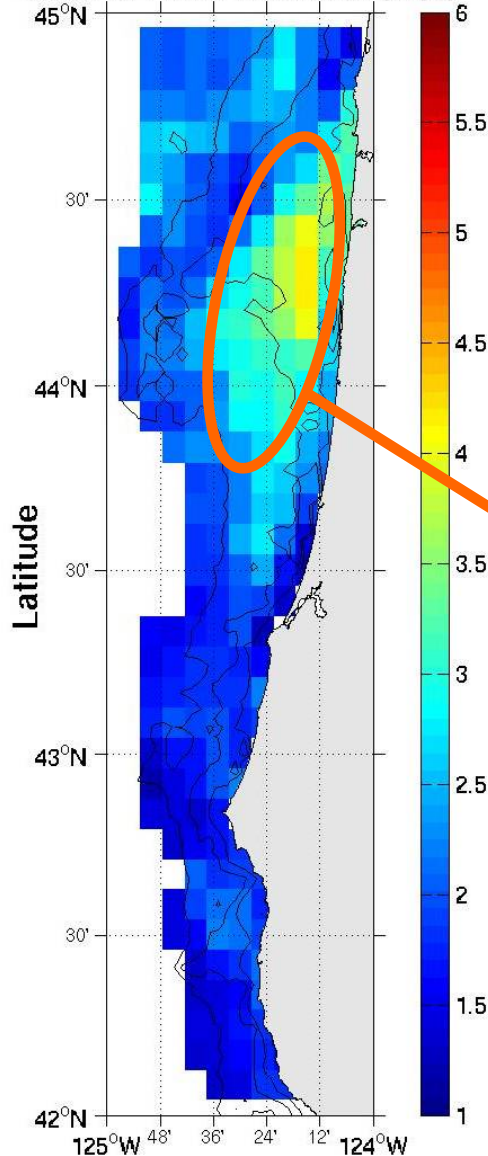
# Comparison of 1 km and 10 km grids

12 Jun 2002

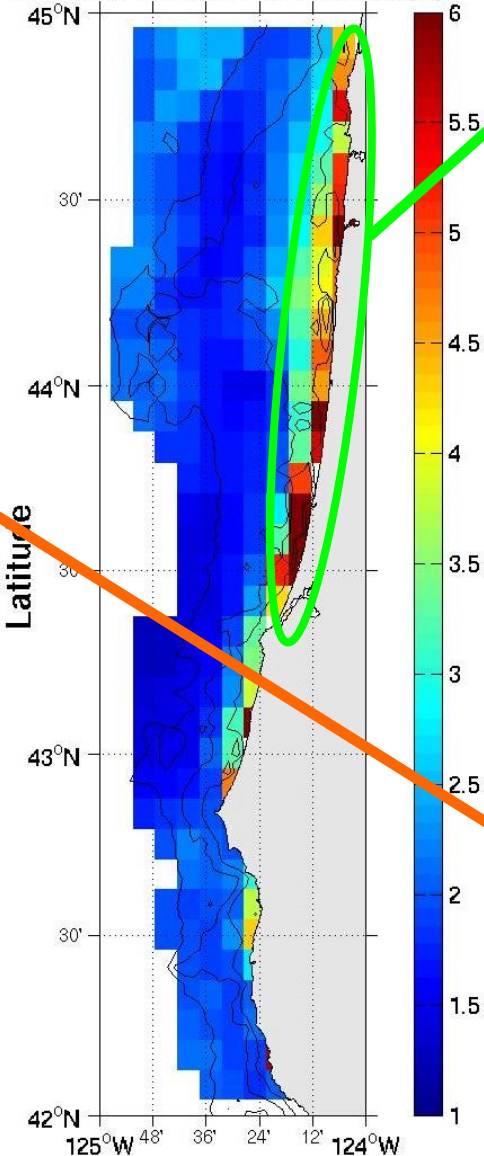


# Comparison of 1 km and 10 km grids

RCCS 2002: Mean e-flush time (days)



NEP4 2002: Mean e-flush time (days)



NEP 10 km simulation shows greatest e-flush times nearshore, likely due to boundary effects

RCCS 1 km simulation shows greatest e-flush times near center of Heceta Bank—not immediately adjacent to shore; does a better job resolving coastal upwelling

# Now for something different...

## Ant Algorithm

- based on the behavior of social insects (like ants, bees)
- social insect interactions are self organized and based on local information
- Ant algorithms have been used to solve large combinatorial optimization problems (e.g., traveling salesman and similar)
- recently used to identify retentive structures in hydrodynamic fields (Segond et al., 2004)

# Ant Algorithm – basics

- Ants move iteratively from one cell to one of 8 neighbors

- direction of stream ( $\alpha$ ); ants are allowed a maximum 45° deviation from stream direction (max of 3 destination cells)

- velocity of the flow in destination cells ( $\mu$ )

- pheromone concentration ( $\phi$ ) in destinations (by population/colony/ant)

- individual ant bias ( $\beta$ )

- my Ant algorithm is derived from, but not identical to Segond et al (2004)

1	2	3
8	A	4
7	6	5

- fitness of 3 destination cells is determined by:

$$F_i = f(\alpha, \beta, \mu, \phi)$$

Scale F so  $\sum F = 1$ ,

Generate UV(0,1) to determine destination



# Ant Assessment of “Potentially” Retentive Regions

Depth = 15 m

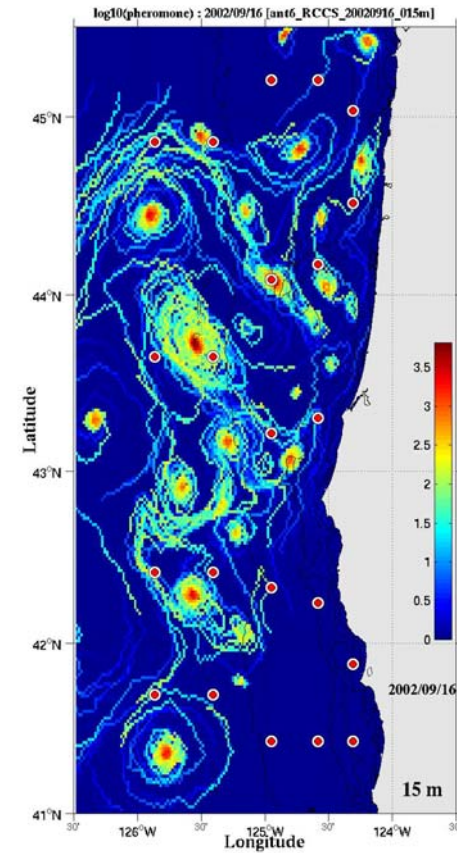
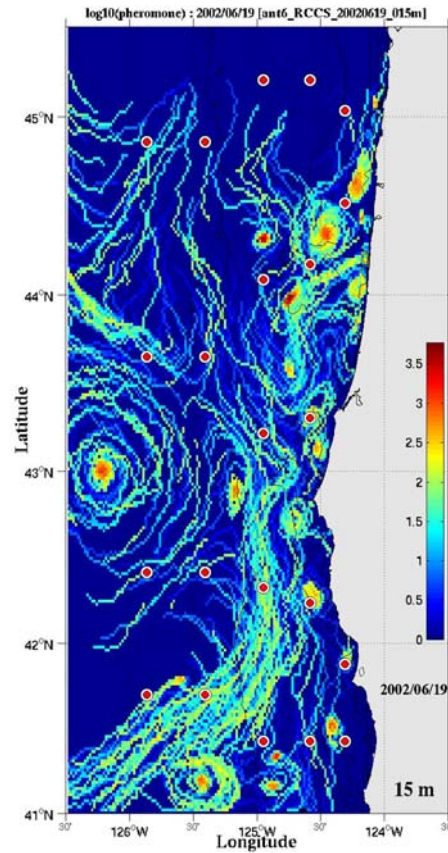
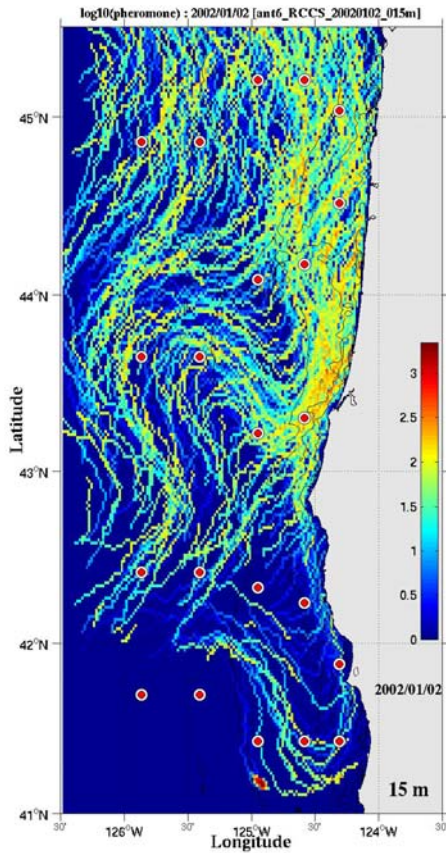
RCCS

22 Initial Colonies of 16 ants/colony

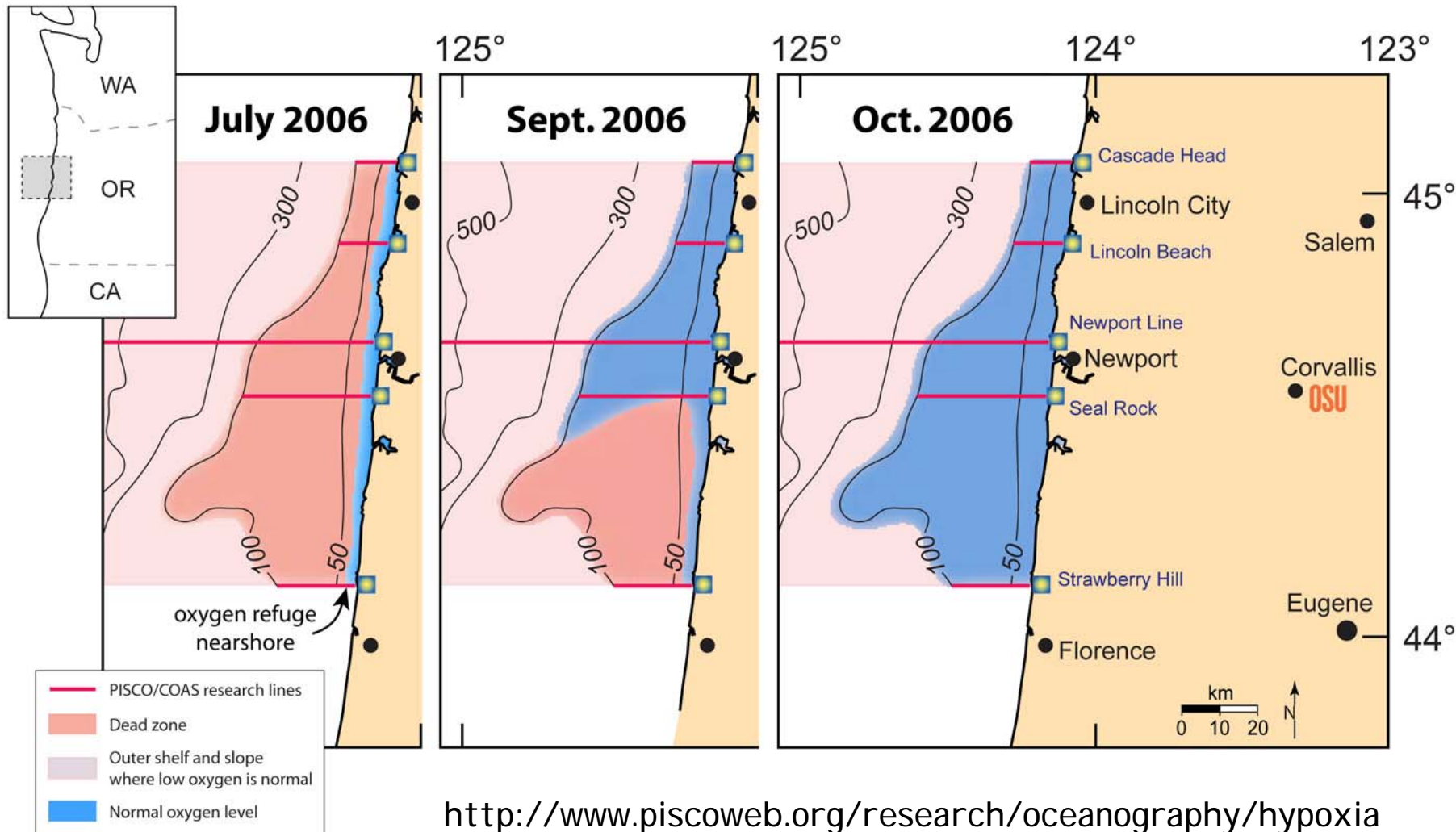
2 Jan

19 Jun

16 Sep



Retention regions matter for productivity and recruitment processes, but also for sink processes. Retention regions may indicate regions susceptible to hypoxia and/or anoxia.



<http://www.piscoweb.org/research/oceanography/hypoxia>

# Summary and Conclusions

- Various metrics for retention and connectivity are available; some have fewer biases than others (boundary encounters in particular can create biases)—e-flushing time has less bias than displacement distance
- Destination maps and source maps are useful for identifying regions that should be high priority for protection (in a MPA sense) and regions that might withstand higher harvest impacts, respectively.
- EOFs of e-flushing time series identify regions of strong co-variability (not shown).
- Retention and connectivity matrices are resolution (scale) dependent (comparison of patterns from 10 km and 1 km resolution models)



# Acknowledgements



- Enrique Curchitser provided the ROMS simulations for which we are very grateful.
- The US National Science Foundation and US National Oceanic and Atmospheric Administration provided funding through the US GLOBEC program that supported this research.

