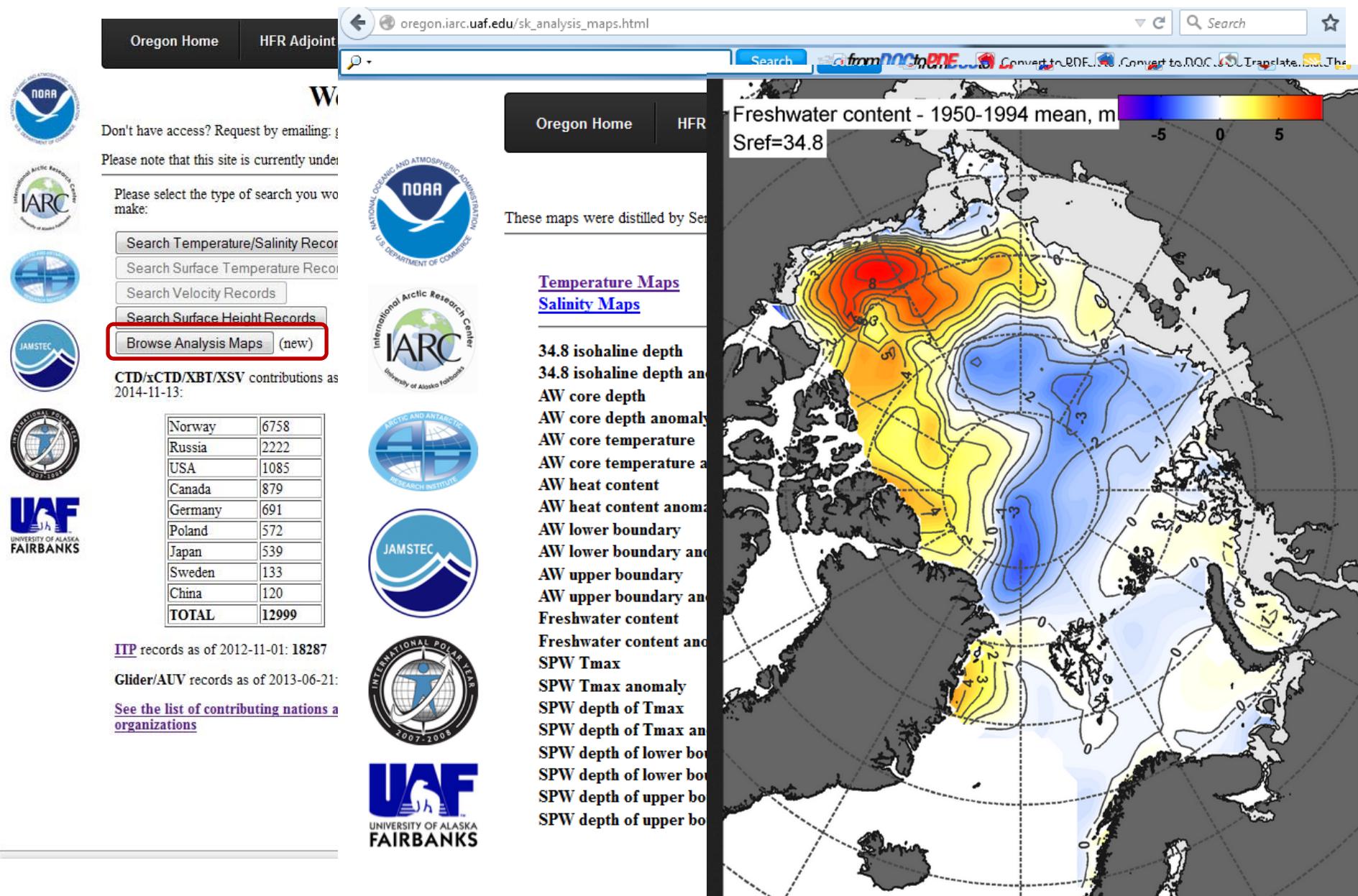


## **Activity during the 8 months**

**G.Panteleev, M.Yaremchuk, J.Stroh, O.Francis,  
Collaboration: P.Stabeno, T.Weingartner, R.Woodgate**

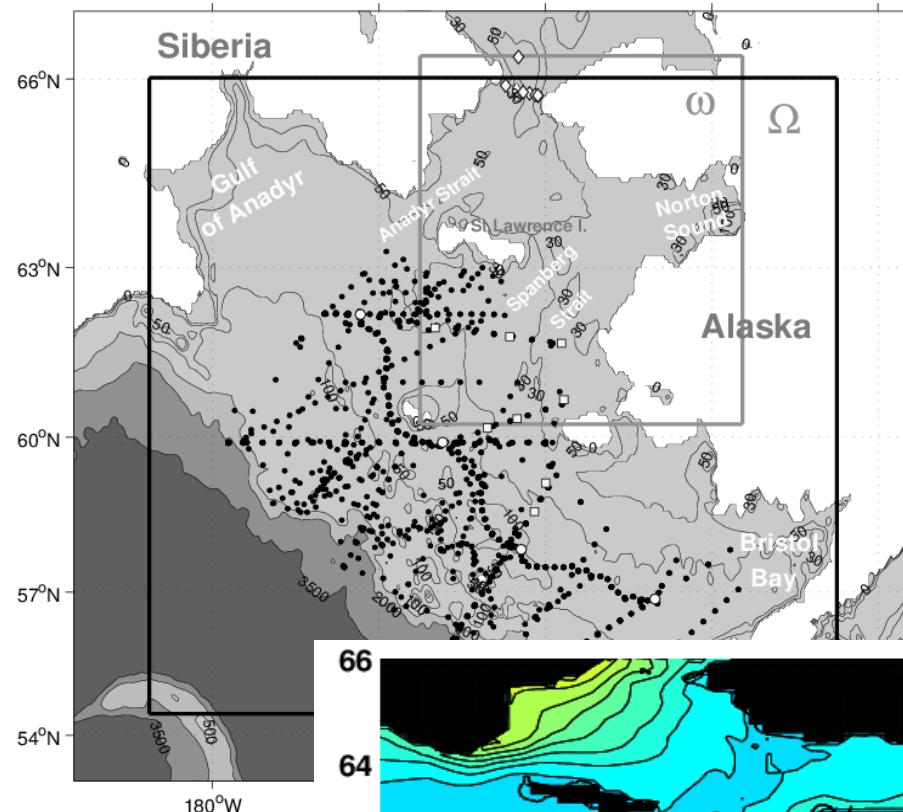
# International Polar Year Database: 2007-2010



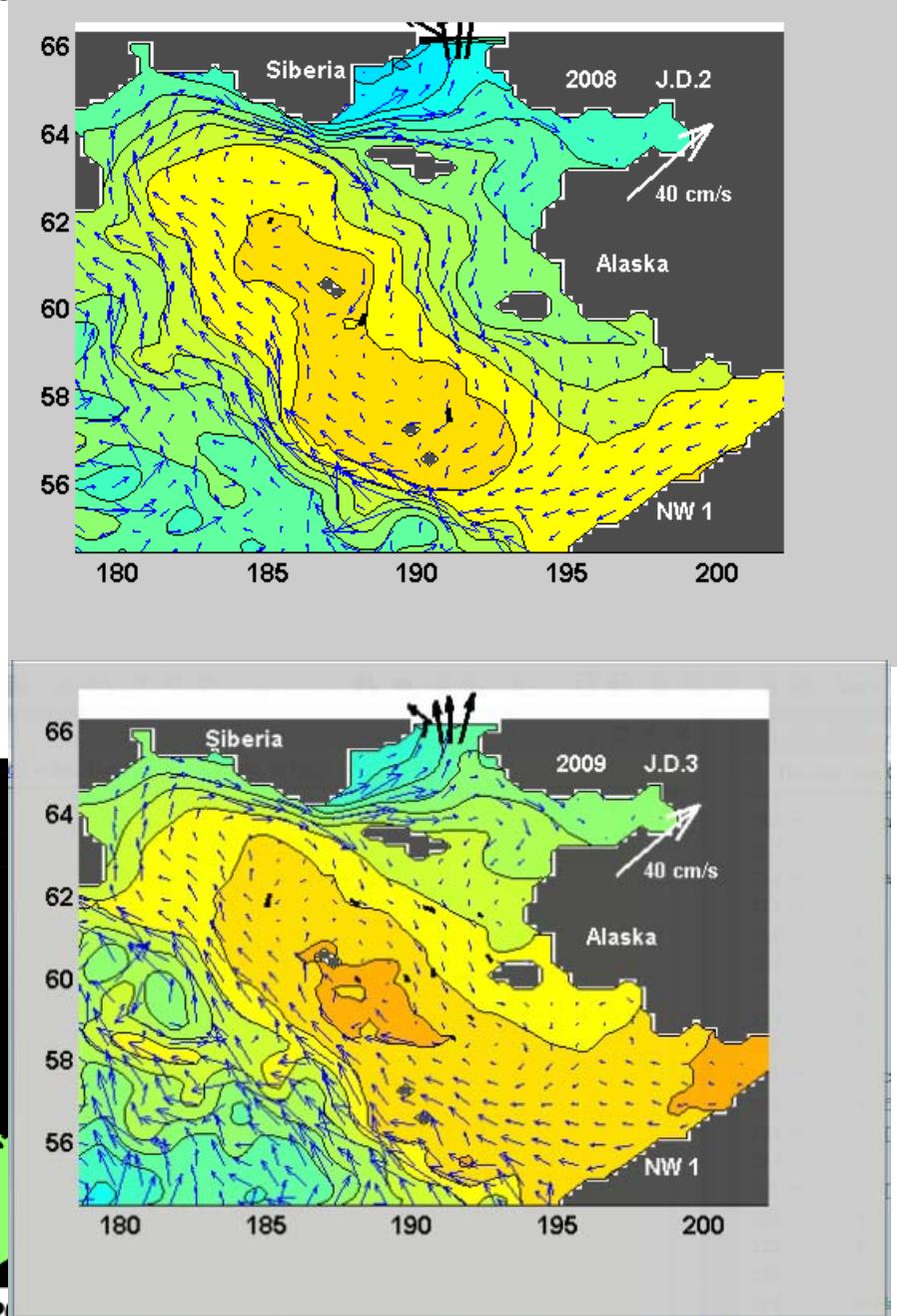
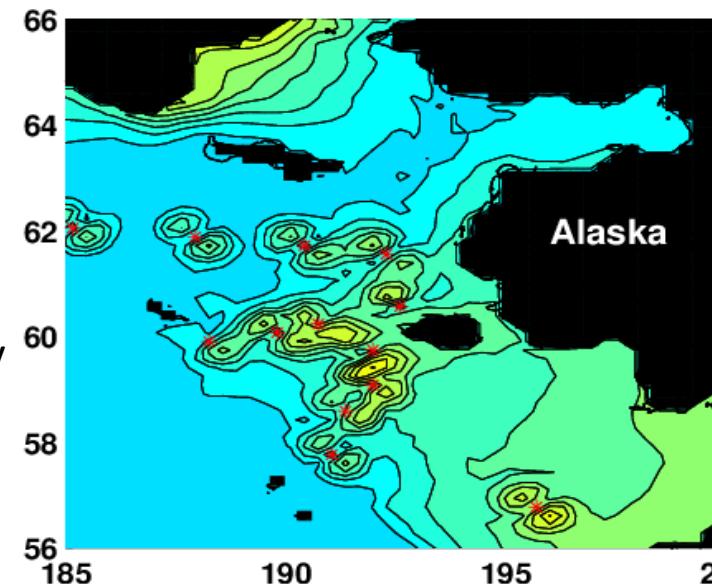
# 4 year reanalysis of the Bering Sea: 2007-2010

Manuscript in preparation: Panteleev, Yaremchuk, Francis, Stabeno

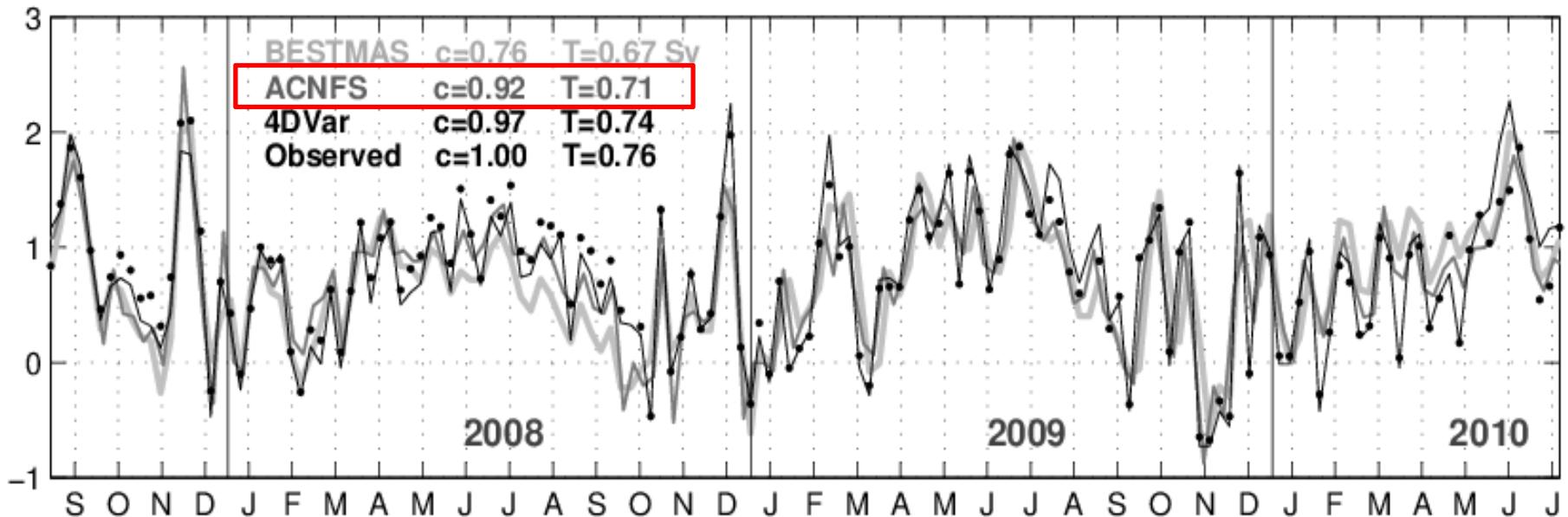
2015.



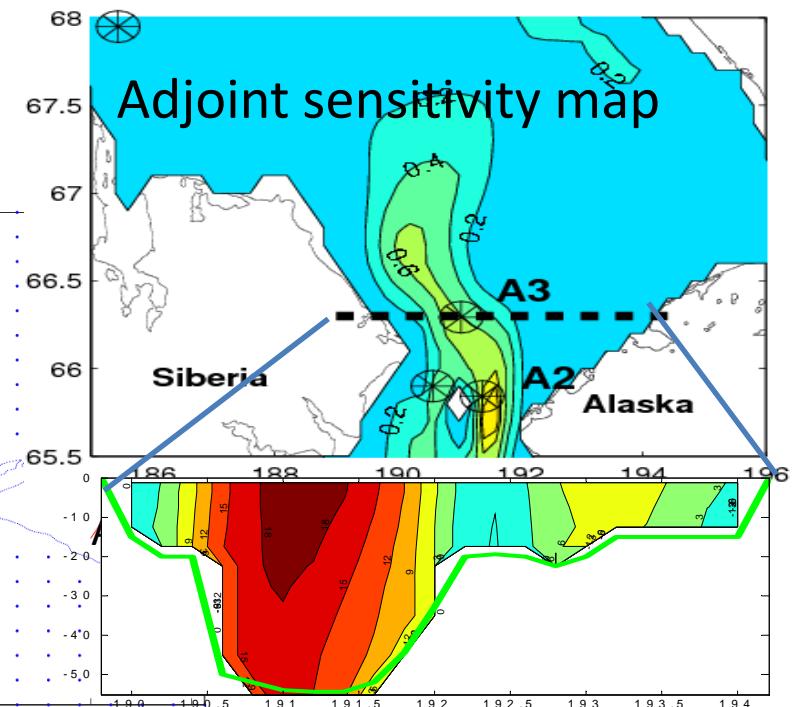
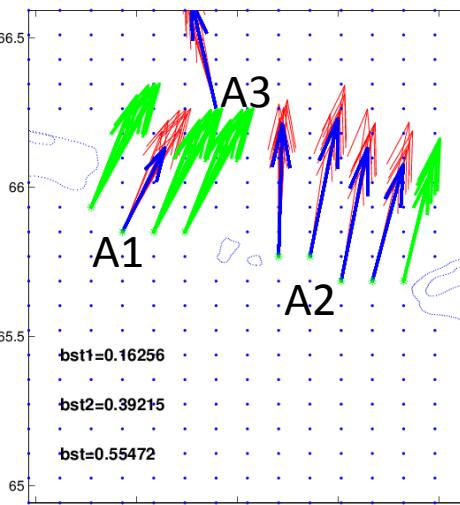
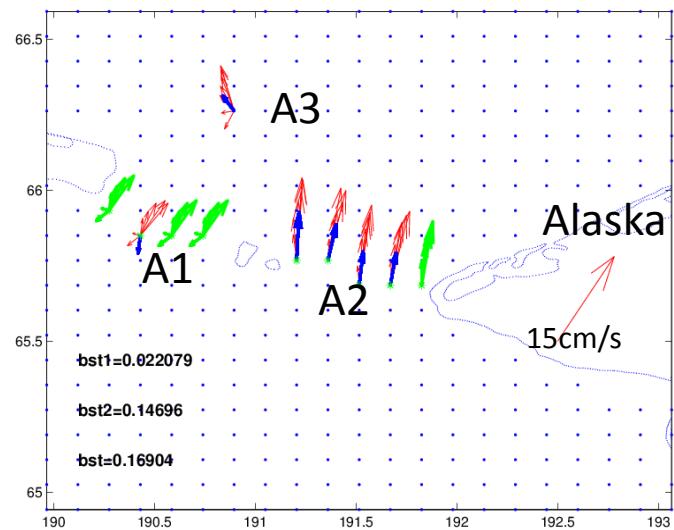
Adjoint  
Sensitivity  
analysis



## Bering Strait transport:

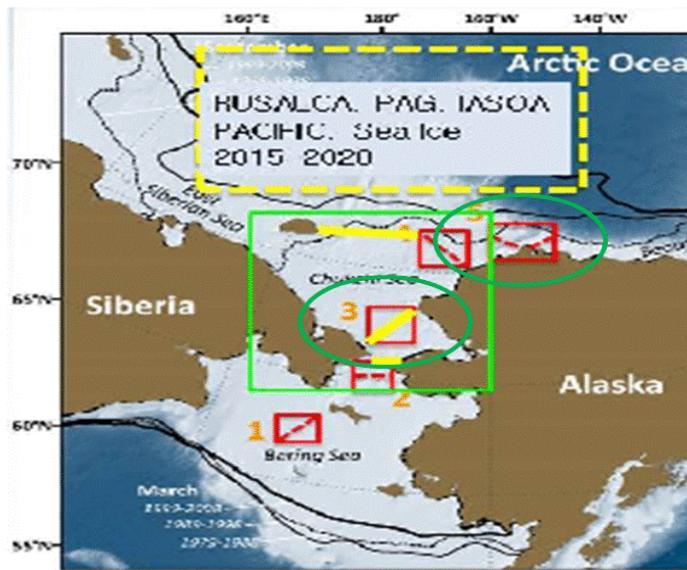


Weekly averaged Correlation( $V_{A3}$ , BST) = 0.96-0.98;  
 Weekly averaged Correlation( $V_{A2}$ , BST) = 0.98-0.99;

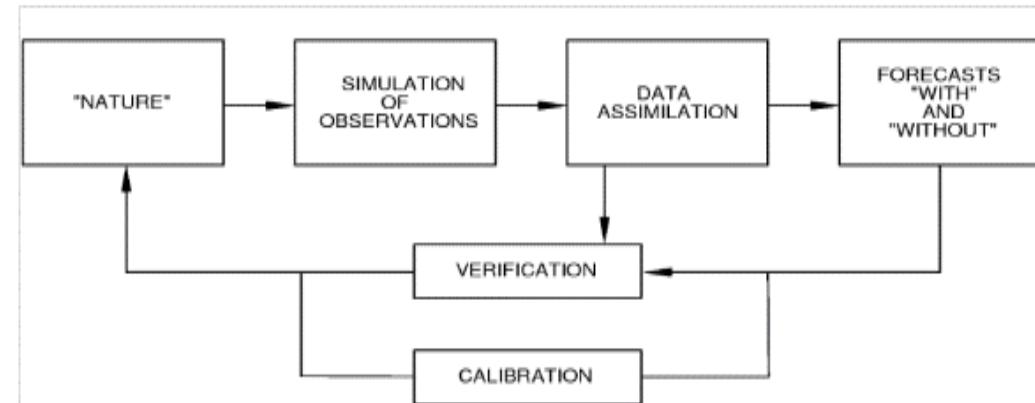


# Passive tracer (biological) sampling

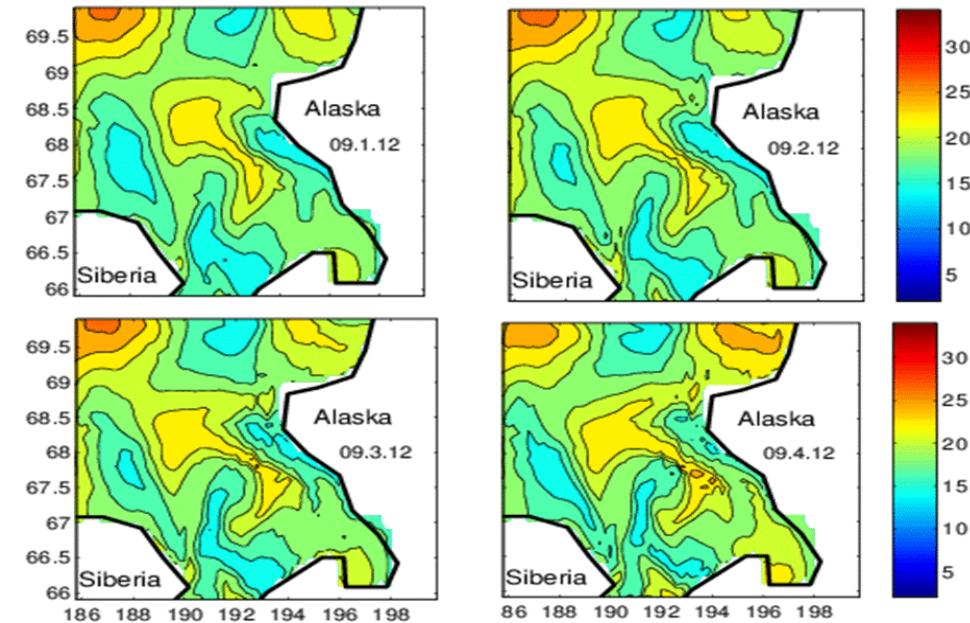
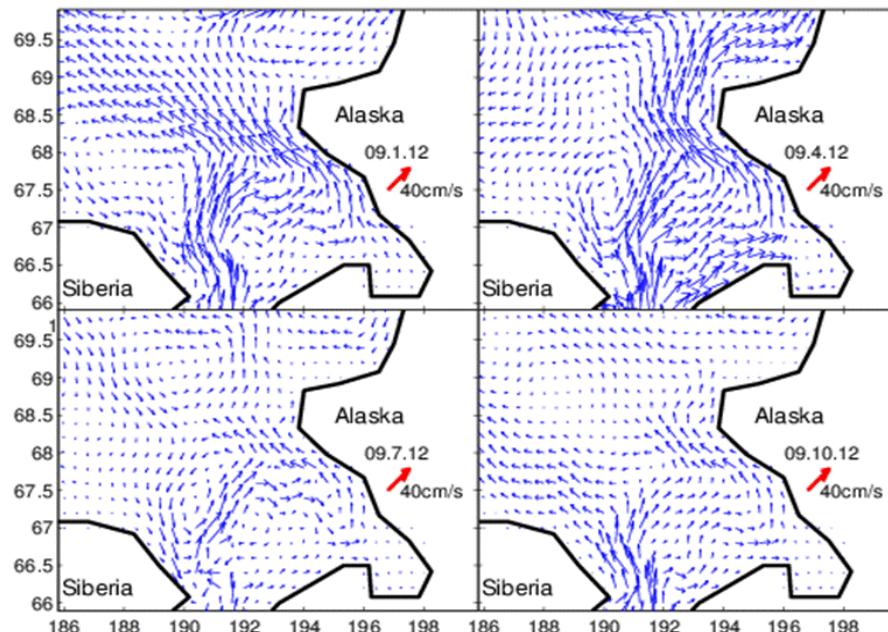
## Area of interest

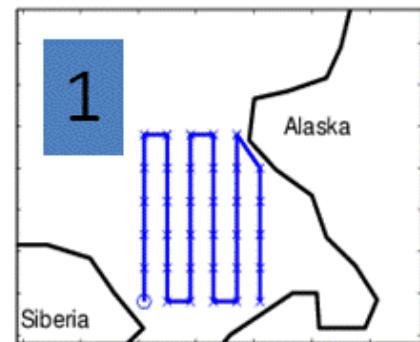
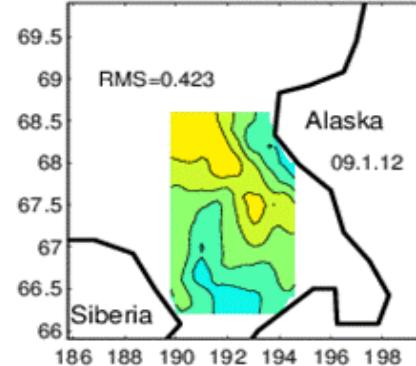
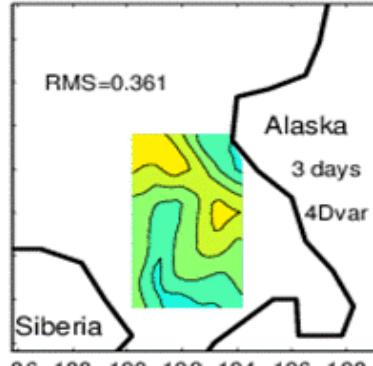
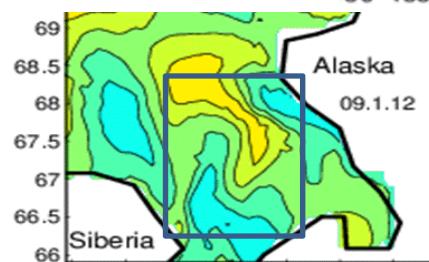
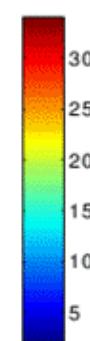
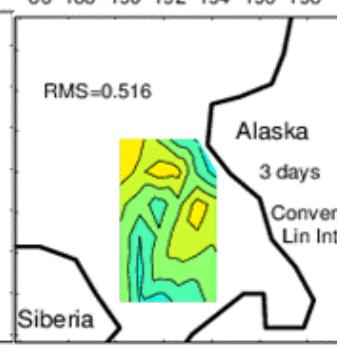
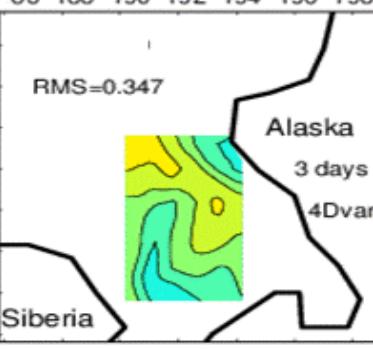
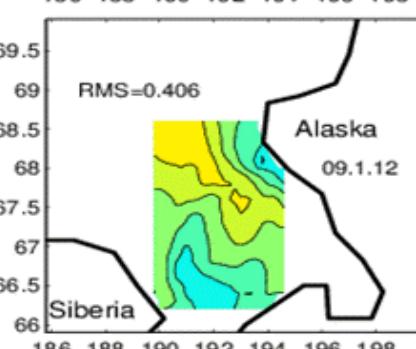
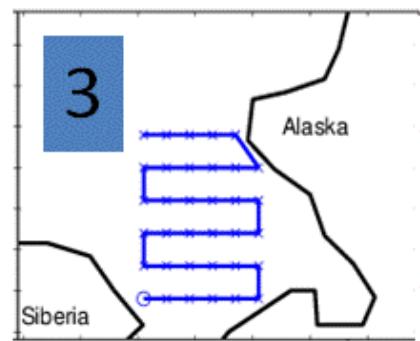
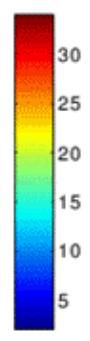
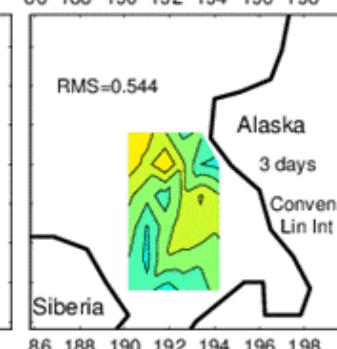
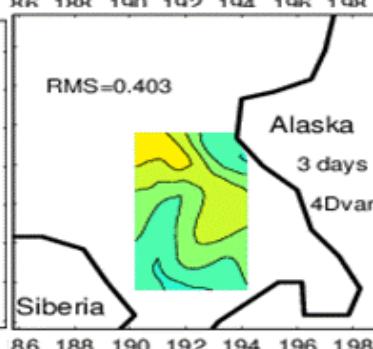
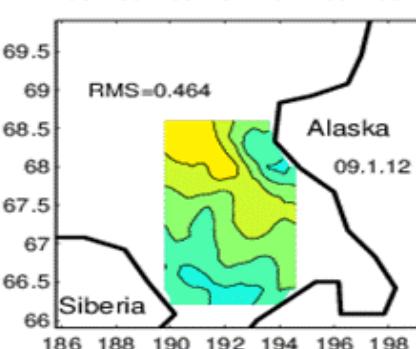
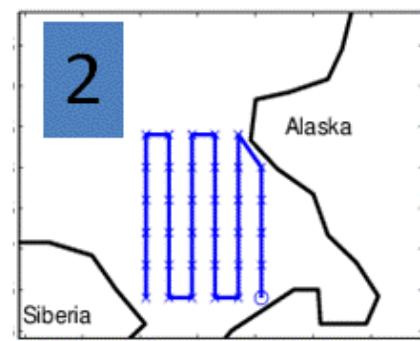
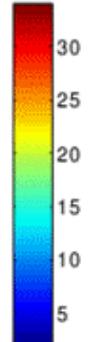
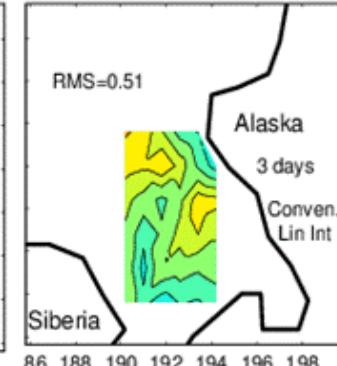


## Observing System Simulation Experiments



Velocity field was derived through the assimilation HFR's and mooring data:



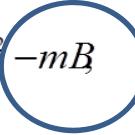
**Surveys****4Dvar Day 1****4Dvar 3days mean****Lin. Interp 3days mean****"TRUE" distribution**

## The 4Dvar advection-diffusion model

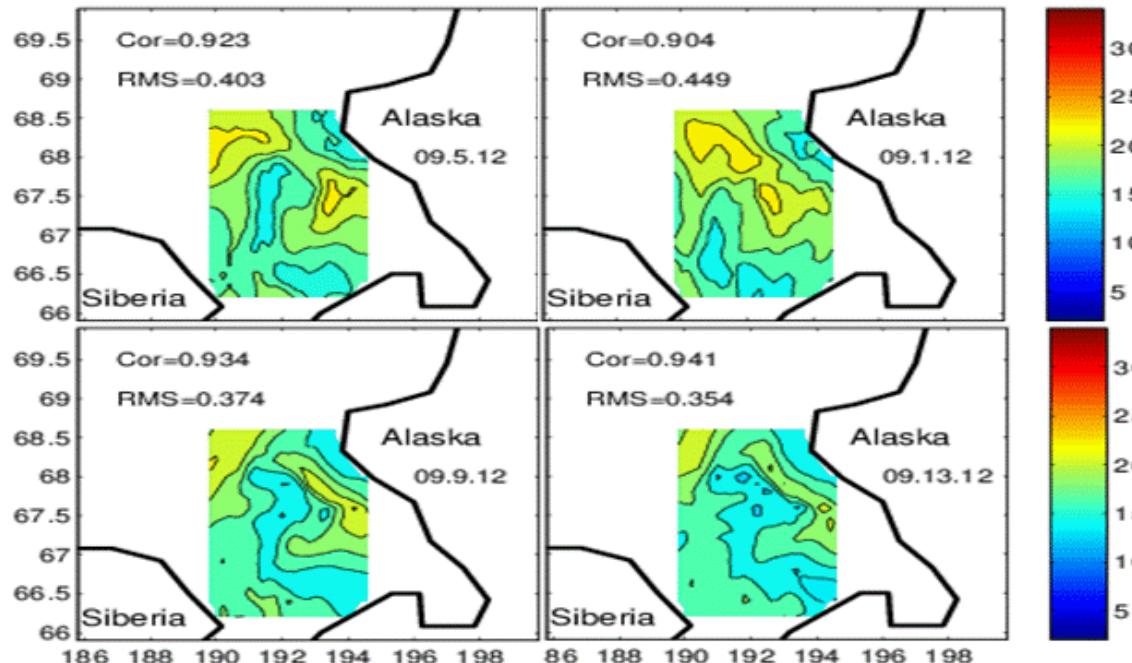
The Advection-Diffusion Biological Tracer Model (ADBTM) is based on modified advection-diffusion equations which are the part of SIOM. Currently ADBTM includes exponential mortality/growth ( $m$ ) and biological tracer horizontal and vertical (sinking) velocity ( $u_B, v_B, w_B$ ) and fluxes  $F_{BO}, F_{BH}$  at the surface and at the bottom:

$$\partial B / \partial t + (u + u_b) \partial B / \partial x + (v + v_b) \partial B / \partial y + (w + w_b) \partial B / \partial z = \Delta B + \partial^2 B / \partial z^2 - mB$$

$$\partial B / \partial z = F_{BO}, z = 0; \quad \partial B / \partial z = F_{BH}, z = H$$

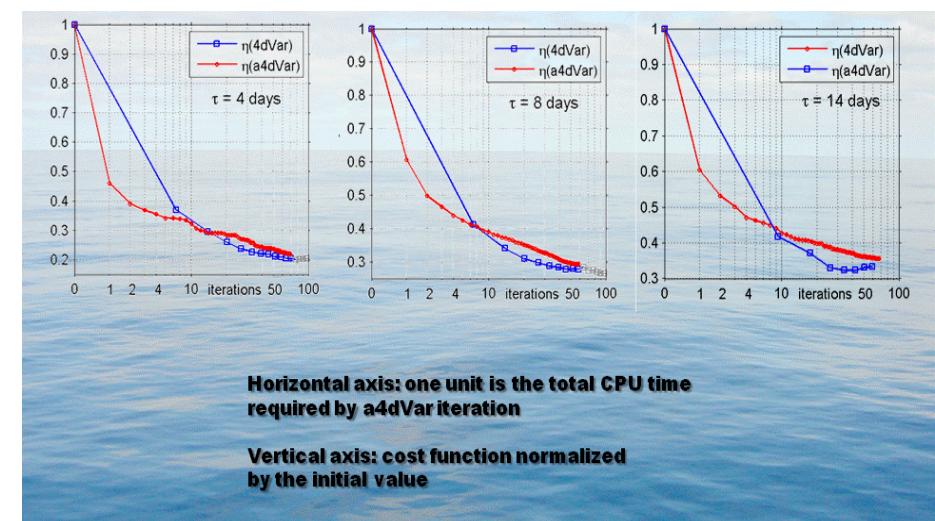
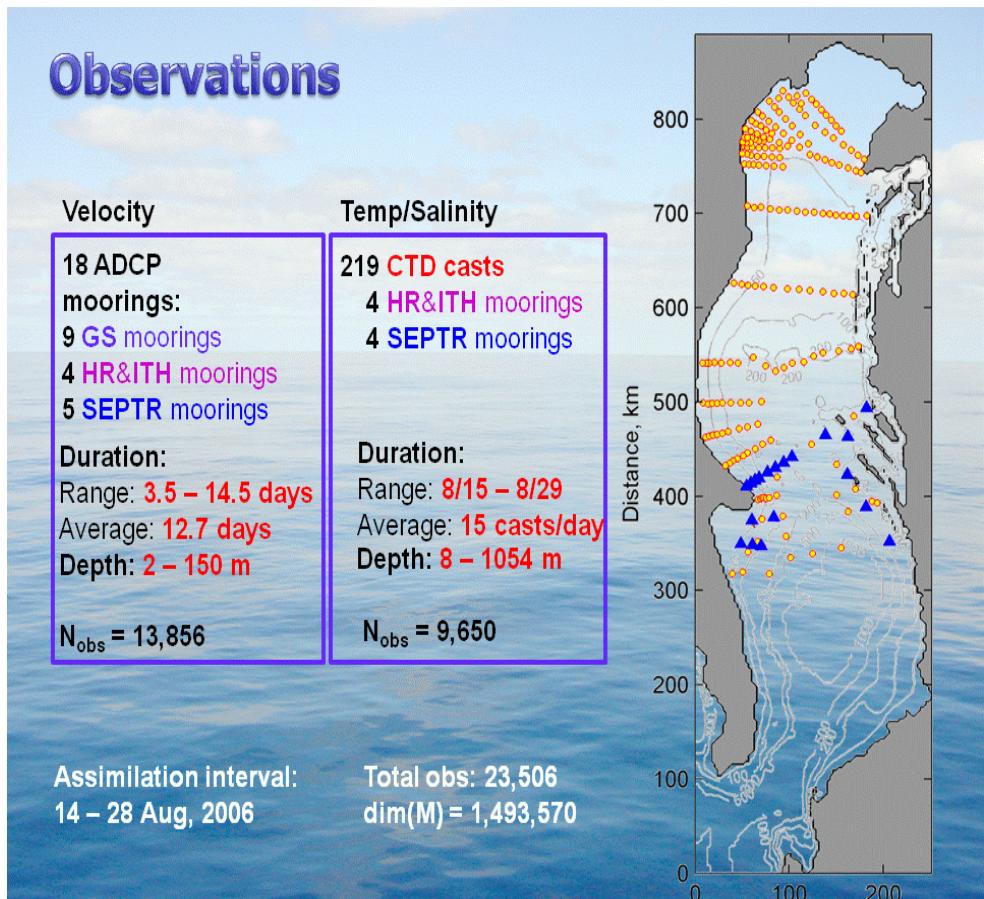
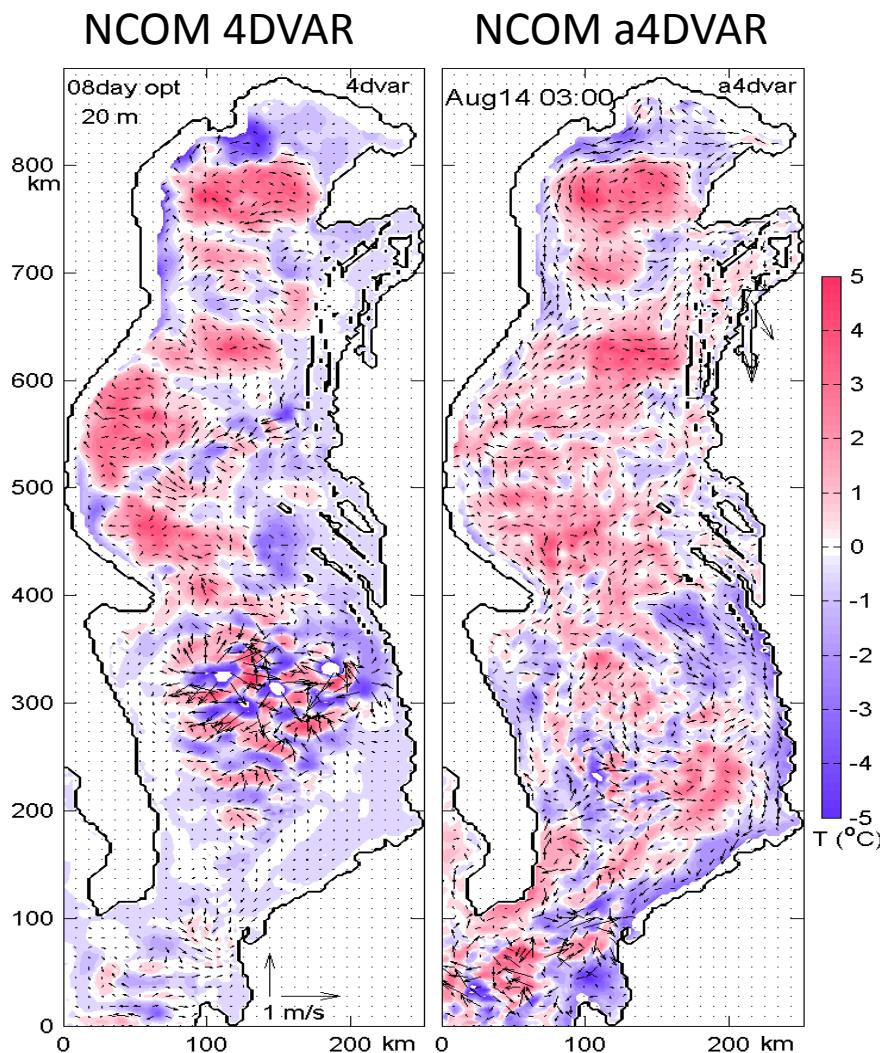


## Estimation of the mortality coefficient



Formally mortality coefficient can be defined from two surveys separated by 1-2 week periods. Survey configuration 1, 3 and 4 were tested with respect of possibility to determine mortality coefficient ( $m$ ) from 2 surveys separated by 7 day period and provided respectively  $m_{optimized} = -1.29e-7, -1.33e-7, \text{ and } -1.4e-7$ . The conventional approach (fitting to exponent) provides  $m_{exp} = -1.53 \text{ -- } -1.57e-7$ . Figure below shows the reconstructed passive tracer evolution with  $m_{optimized} = -1.29e-7$  for the survey configuration #1.

# Adjointless 4Dvar NCOM: extention to the ICE-HYCOM and ICE-ROMS



## **Conclusions:**

1. SAON/NOAA International data base provide the one stop for CTD observations in the AO during 2008-2010. There is a strong need to continue this activity and develop PAG database.
2. 2007-2010 Northern Bering Sea reanalysis has been accomplished. The lessons from this study are:
  - a) preliminary adjoint sensitivity analysis and OSSE help to increase the information content of the observation. That will result to the more accurate estimates of the ocean state
  - b) HYCOM ACNFS (3Dvar) provides very good estimates of the Bering Strait transport and near surface circulation. It can be used as a first guess for the incoming Adjointless ROMS-HYCOM DAS'es
3. A4Dvar NCOM has been tested. A4Dvar ROMS-ICE, HYCOM-ICE is the next step.
4. Variational inversion of the biological observations using the passive traces allows to increase accuracy of the estimates of the biological parameters and mortality coefficient.