TESTING TOXIN PREDICTION MODELS IN A CALIFORNIA HAB HOTSPOT

<u>Clarissa Anderson^{1,2}</u>*, Raphael Kudela¹, Claudia Benitez-Nelson², Emily Sekula-Wood², Yi Chao³, Thomas Ellis⁴, David Siegel⁵, Libe Washburn⁵, Gregg Langlois⁶

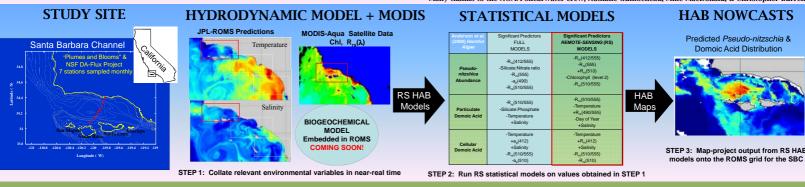
> ¹UC Santa Cruz, ²University of South Carolina, ³NASA Jet Propulsion Laboratory-UCLA, ⁴ NASA Airborne Sensor Facility, ⁵UC Santa Barbara, ⁶CA Department of Public Health **corresponding author: clrander@ucsc.edu*

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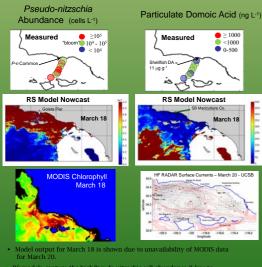
ABSTRACT

The documented link between upwelling-related physical signatures, macronutrients, and toxigenic *Pseudo-nitzschia* blooms in the various "hotspots" throughout California has motivated attempts to forecast these harmful algal blooms (HABs) as a function of select environmental variables. The Santa Barbara Channel (SBC), CA is one such localized region for toxic dilatom blooms in the California Current System with nearly annual toxic events occurring in spring and summer months since 2002. Existing statistical models that use physical, chemical, and bio-optical observations for predicting *Pseudo-nitzschia* abundance and domoic acid (DA) concentration in the SBC are currently being tested in a nowcast mode using spatially-explicit predictions of environmental variables from MODIS ocean color geophysical data and a Regional Ocean Modeling System (ROMS) model for the SBC. Relevant thresholds for each model generate "event" forecast, and results are validated relative to on-going monthly observations along a channel-wide transect to assess the potential use of model precise physical physical physical physical currents. Additional comparisons of model output with domoic acid levels in commercial mussels collected by the CA Department of Public Health demonstrate the disconnect in the relationship between in situ bloom dynamics and the toxin loads measured in nearshore shellfish beds.

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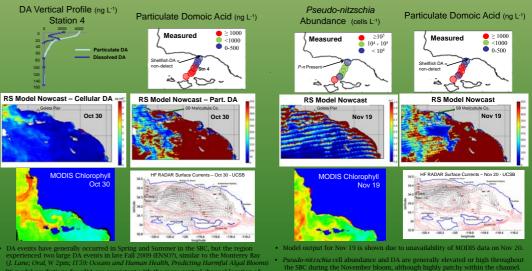


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- RS models capture the high *Pseudo-nitrschia* cell abundance & low to moderate pDA measured across the SBC; low nearshore pDA is reflected in the low shellfish DA levels collected from the SBMC site (regulatory limit = 20 µg g⁻¹).
- Elevated DA at mid-channel stations is not reflected in the DA model, but may be a resul of the mesoscale eddy seen in CODAR surface current structure and possibly in the MODIS Chi image for March 18 when an eddy feature was already spinning up.
- Cyclonic, convergent eddies in the SBC are often associated with high productivity over the basin (Brzezinski & Washburn, *in review*) and development of a local hotspot of DA during HAB events (Anderson et al., 2006, MEPS).

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RS model predictions for pDA are consistent with the more central-channel location of high observed DA concentrations, indicating that the model captures broad patterns in DA.

 Elevated cellular DA levels mid-channel are roughly inverse of pDA maxima and may be due to the presence of a cyclonic eddy (CODAR) that appeared to be concentrating biomass (MODIS) over the central to vestern basin where the <u>vertical profile</u> of very high subsurface pDA & <u>dissolved DA</u> indicates sinking of a late phase surface bloom or the presence of a subsurface bloom maximum.

 DA in shellfish was below detection with Jellett kits from Aug-Dec 2009, but DArelated sea lion strandings were reported from SF to the SBC from Sep-Dec 2009.

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tially retentive, cyclonic eddy (CODAR) and high pDA/dDA down to 30 m

This spatial heterogeneity appears in the RS Model predictions of cell and DA, however satellite noise artifacts potentially increase patching

filled with the DINEOF interpolation program to improve temporal resand overall lead times of HAB predictions.