Tracking, Understanding and Predicting Toxic Phytoplankton blooms and their effects on King Scallops populations in the Bay of Seine (Task 5: ANR Systerra-COMANCHE)



M. Schapira¹, R. Le Gendre¹, L. Fiant¹, J. Fauchot², V. Raimbault², C. Dreanno³, P. Claquin², P. Riou¹

¹IFREMER, Laboratoire Environnement et Ressource de Normandie, Avenue du Général de Gaulle, BP 32, 14520 Port en Bessin, France. ²Equipe Biologie des Ecosystèmes Côtiers, UMR 100 IFREMER - PE2M, Université de Caen Basse-Normandie, Esplanade de la Paix 14032 Caen, France.³IFREMER, Service Interfaces et Capteurs, Unité Recherches et Développements technologiques, Technopôle, 29280 Plouzané, France.

BAY OF SEINE: 2 MAJOR TOXIC EVENTS

2004 – Amnesic Shellfish Poisoning (ASP)



- [DA] > sanitary threshold (i.e. >20 µg DA g⁻¹ ww) from Nov. 2004 to Jan. 2005 (Fig. 1B)

2 *Pseudo-nitzschia* species were identified as the potential source of DA during the 2004 toxic event (Nézan et al. 2006)



✤ King scallops, Pecten *maximus,* is the first species in landing (in tons and value) for the fishing fleet in the Bay of Seine.

Dinophysis sp. was identified as the source of OA during the 2005 toxic event (Amzil et al. 2007)

2005 – Diarrheic Shellfish Poisoning (DSP)



AN IMPROVED UNDERSTANDING OF THE DETERMINISM OF THESE TOXIC BLOOMS IS CRITICALLY NEEDED TO DEVELOP LONG TERM MANAGEMENT STRATEGIES FOR KING SCALLOPS FISHERIES IN THE BAY OF SEINE



UNDERSTANDING PAST EVENTS & MONITORING TOXIC PHYTOPLANKTON BLOOMS



Survey ASP and DSP toxicity levels in king scallops in relation with the proliferation of toxic phytoplankton blooms (French Phytoplankton and Phycotoxin Monitoring Network, REPHY)

Study past toxic events (ASP & DSP) in relation with the variability of environmental parameters and climatic events (REPHY and RHLN data; Normandy Hydrology Monitoring Network)



Fig 3: Standardized abundances of *Pseudo-nitzschia* sp. recorded at the REPHY sampling site 'Cabourg' (cf. Fig. 5) from 2002 to 2010 (data source, REPHY). The green arrow indicates the 2004 toxic event. The red line represents the sanitary threshold (i.e. 100 000 cell I-1).

LIMITATIONS

Identification of toxic species Eco-physiology poorly documented

Fig 4: Standardized abundances of *Dinophysis* sp. Recorded at the REPHY sampling site 'Antifer' (cf. Fig. 5) from 2002 to 2010 (data source, REPHY). The green arrow indicate the 2005 toxic event. The black line represents the sanitary threshold (i.e. 500 cells l-1).



microarray been * A has designed using oligoprobes (25 mers) matched to toxic microalgae ribosomal RNA.

Pseudo-nitzschia sp. * For labelled target DNA was prepared polymerase chain reaction by amplification of ITS region using a Cy5-labeled primers. DNA was extracted from monoclonal cultures. Hybridization performed was according to the method described by Le Berre et al. (2003).

Pau 6-6)



• **Preliminary data** show that the current chips can specifically detect and discriminate P. americana, P. pungens, P. australis, P. multiseries and P. fraudulenta.

ECOPHYSIOLOGY OF TOXIC PHYTOPLANKTON



✤ Identification of the different Pseudo-nitzschia strains present in the Bay of Seine using Transmission Electronic Microscopy (TEM).

Pseudo-nitzschia sp. TEM



ANTIFER

- ✤ Isolation & culture of the different Pseudo-nitzschia strains
- **Eco-physiology** of the different *Pseudo-nitzschia* strains



Temperature (°C) From Claquin *et al*. 2008

P. Fraudulenta

• Biological & physiological "ID card" for each strains isolated in the Bay of Seine





PREDICTING TOXIC PHYTOPLANKTON BLOOMS IN THE BAY OF SEINE

The Hydro-biological model, MARS3D has recently been refined for the Bay of Seine (500 m resolution).

Different phytoplankton groups (i.e. diatoms, dinoflagellates and nanophytoplankton) have been incorporated to the physical model (cf. Fig. 1&2).

This model has been validated with the data set provided by the RHLN (Normandy Hydrology Monitoring Network).

Finally a representation of a toxic diatom - *Pseudo-nitzschia* –will be added.

References cited: Amzil et al. (2007) Sixth International Conference on Molluscan Shellfish Safety, 307-314. Claquin et al. (2008) Aquatic Microbial Ecology, 51, 1-11. Le Berre et al. (2003) Nucleic acid Research, 31 (16) e88. Nézan et al. (2006) Harmful Algae News, 31, 1-3.

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