## Tracking, Understanding and Predicting

# Toxic Phytoplankton blooms and their effects on King Scallops populations

in the Bay of Seine

(Task 5: ANR Systerra-COMANCHE)

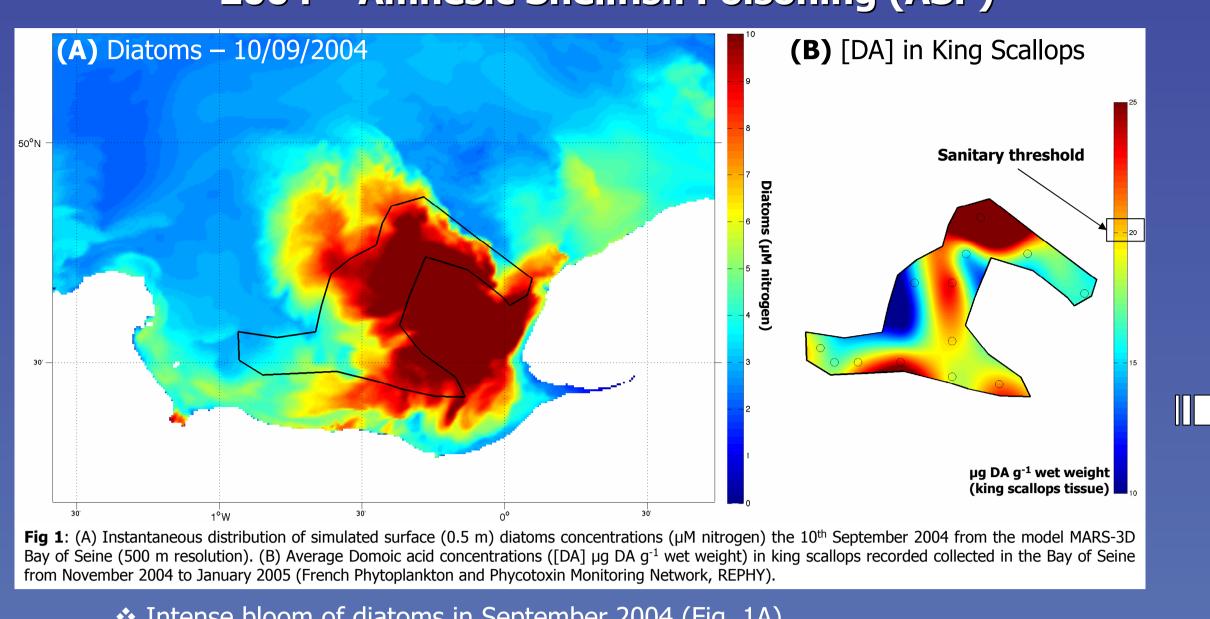


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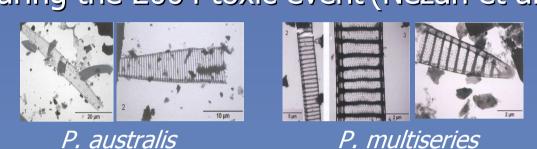
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#### **BAY OF SEINE: 2 MAJOR TOXIC EVENTS**

#### 2004 – Amnesic Shellfish Poisoning (ASP)



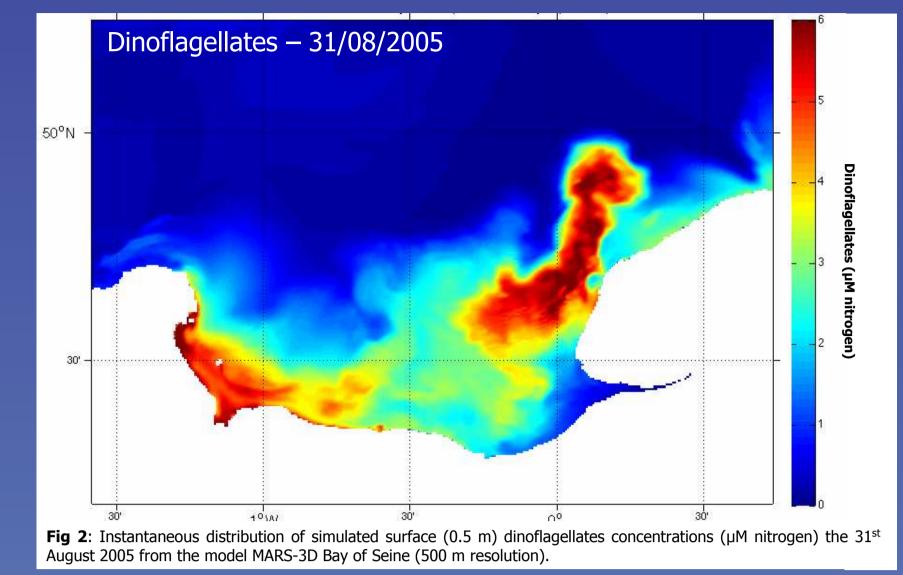
- ❖ Intense bloom of diatoms in September 2004 (Fig. 1A)
- [DA] > sanitary threshold (i.e. >20 µg DA g<sup>-1</sup> ww) from Nov. 2004 to Jan. 2005 (Fig. 1B)
  - 2 Pseudo-nitzschia species were identified as the potential source of DA during the 2004 toxic event (Nézan et al. 2006)



**CLOSURE** OF KING SCALLOPS **HARVESTING SITES** IN 2004 & 2005

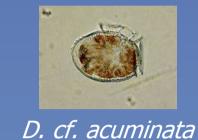
❖ King scallops, Pecten maximus, is the first species in landing (in tons and value) for the fishing fleet in the Bay of Seine.

#### 2005 — Diarrheic Shellfish Poisoning (DSP)



- ❖ Many blooms of dinoflagellates were observed in July/August 2005 (Fig. 2)
- ❖ Positive DSP mouse bioassays & very high [OA+DTX3] from Oct. to Dec. 2005 (Amzil et al. 2007)

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

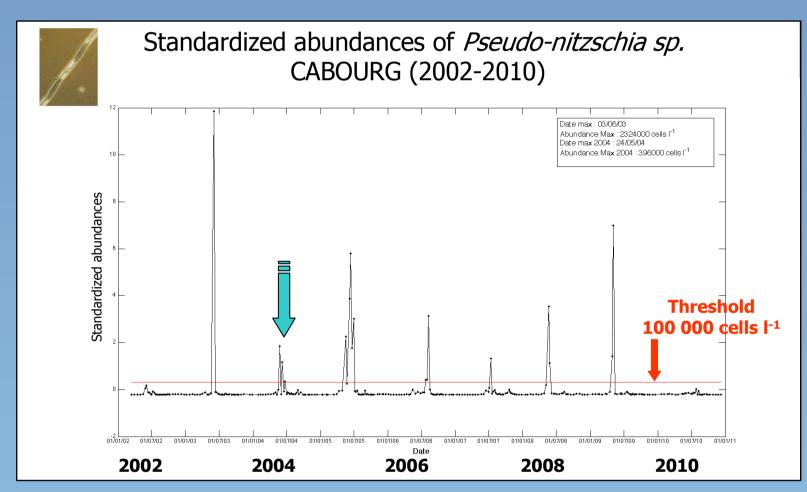


AN IMPROVED UNDERSTANDING OF THE DETERMINISM OF THESE TOXIC BLOOMS IS CRITICALLY NEEDED TO DEVELOP LONG TERM MANAGEMENT STRATEGIES FOR KING SCALLOPS FISHERIES IN THE BAY OF SEINE

ONE TASK OF THE COMANCHE PROJECT

(Ecosystems interactions and anthropogenic impacts on King scallops populations in the English Channel)

## UNDERSTANDING PAST EVENTS & MONITORING TOXIC PHYTOPLANKTON BLOOMS



**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 l-1).

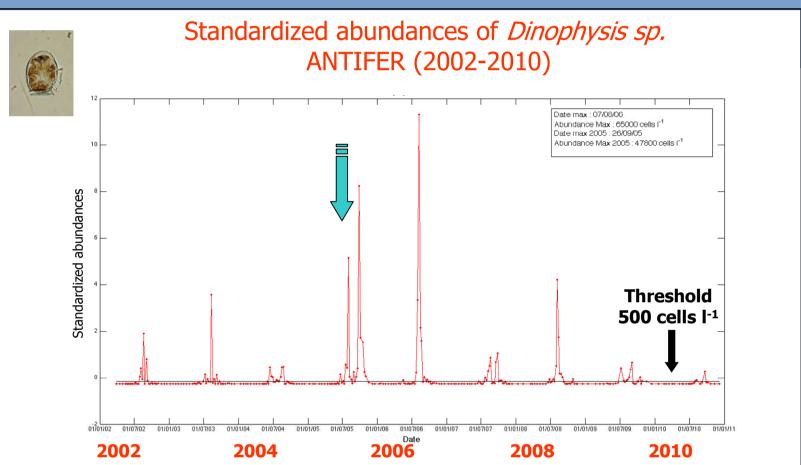
- Survey ASP and DSP toxicity levels in king scallops in relation with the proliferation of toxic phytoplankton blooms (French Phytoplankton and Phycotoxin Monitoring Network, REPHY)
- ❖ Study past toxic events (ASP & DSP) in relation with the variability of environmental parameters and climatic events (REPHY and RHLN data; Normandy Hydrology Monitoring Network)



Identification of toxic species

Absent

Eco-physiology poorly documented

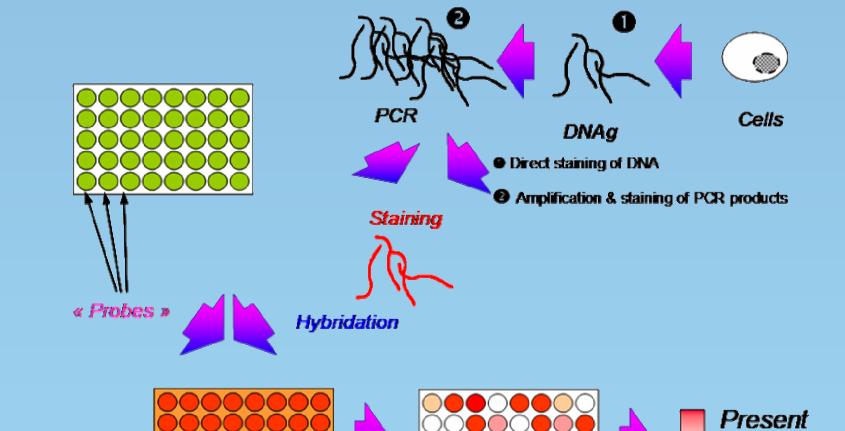


## **CABOURG** Fig 5: Location of the REPHY sampling sites 'Cabourg' and 'Antifer. **Fig 4**: Standardized abundances of *Dinophysis* sp. Recorded at the REPHY sampling site 'Antifer' (cf. Fig. 5) from 2002 to 2010 (data source, REPHY). The green arrow indicate the 2005 toxic event. The black line represents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

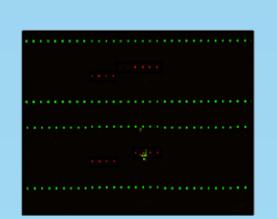
## TRACKING:

### DEVELOPMENT OF DNA MICROARRAYS FOR PHYTOPLANKTON IDENTIFICATION = "PHYTOCHIPS"

- microarray has designed using oligoprobes (25 mers) matched to toxic microalgae ribosomal RNA.
- Pseudo-nitzschia sp, labelled target DNA was prepared polymerase chain reaction amplification of ITS region using a Cy5-labeled primers. DNA was extracted from monoclonal cultures. Hybridization was performed according to the method described by Le Berre et al. (2003).



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



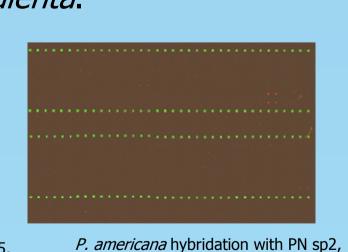
(Pau1-4, Pau5-1, Pau5-2, Pau5-3,

Pau6-1, Pau 6-2, Pau6-4, Pau 6-5 et

Pau 6-6)

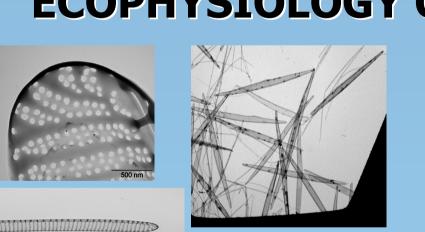
P. pungens: 7 specific probes (1-5, 2-2, 3-5,

4-6, 5-1, 6-3, 7-2)



Americana2, Americana4

## **ECOPHYSIOLOGY OF TOXIC PHYTOPLANKTON**



Light

**♦** T℃

Nutrients

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

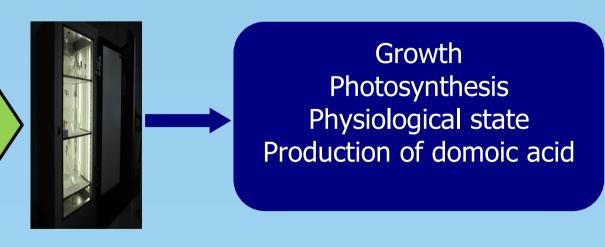


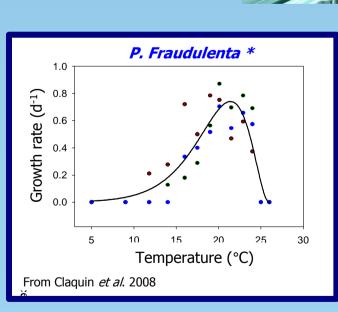
Bay of Seine

ANTIFER

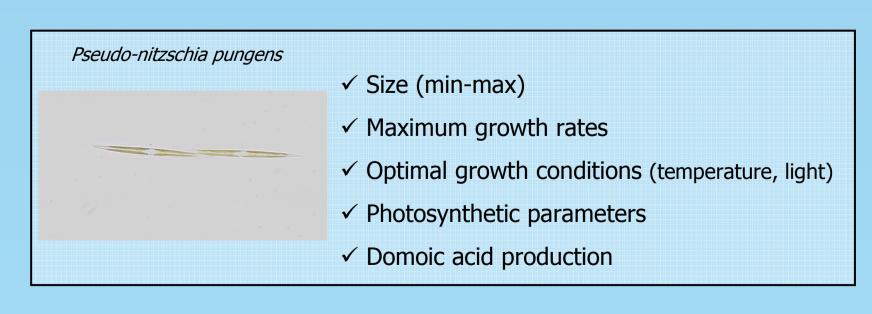
- ❖ Isolation & culture of the different *Pseudo-nitzschia* strains
- **Eco-physiology** of the different *Pseudo-nitzschia* strains

Pseudo-nitzschia sp. TEM





❖ 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.



Nézan et al. (2006) Harmful Algae News, 31, 1-3.