

Phytochip: a new tool to study the diversity of toxic phytoplankton in the Bay of Seine

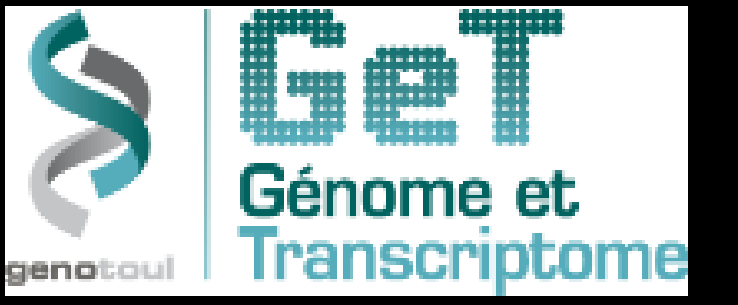
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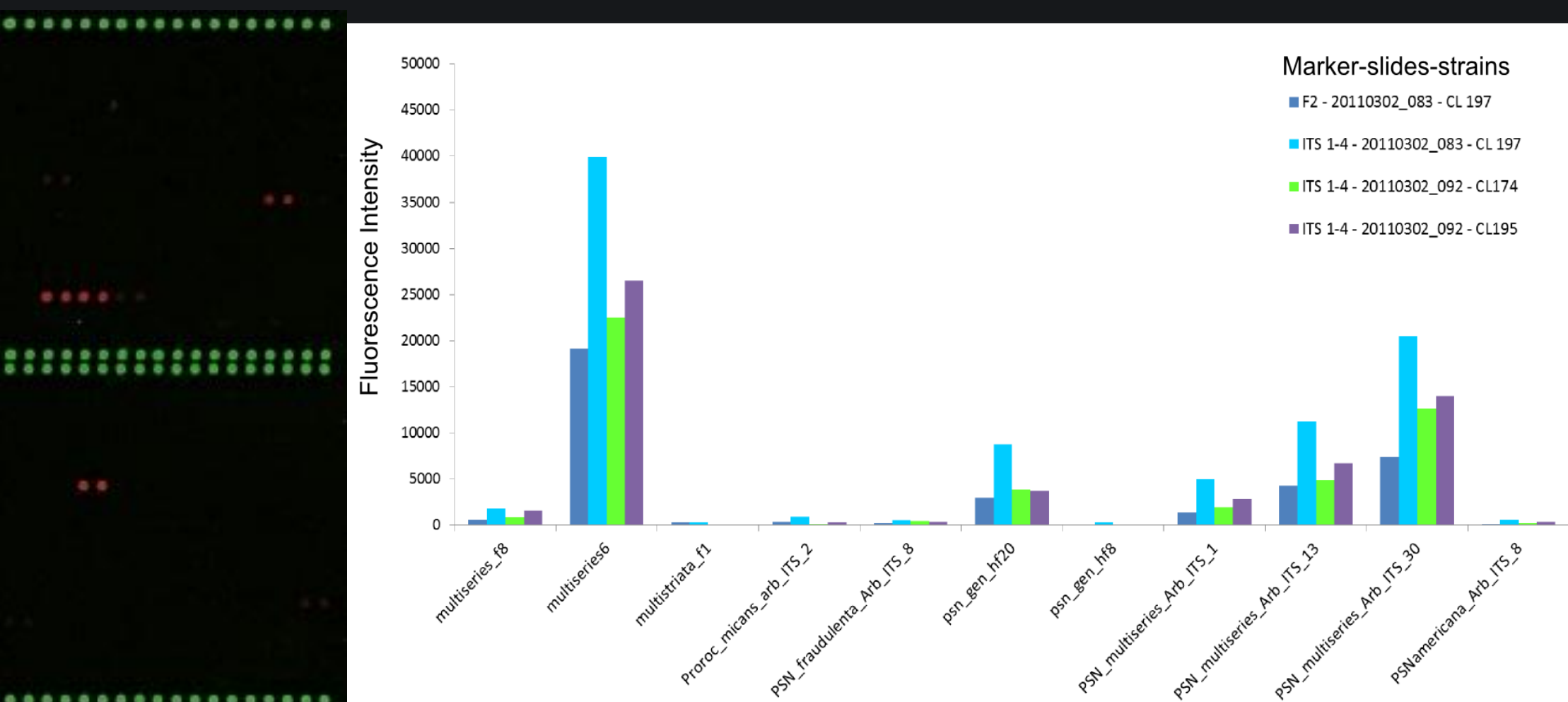


Introduction

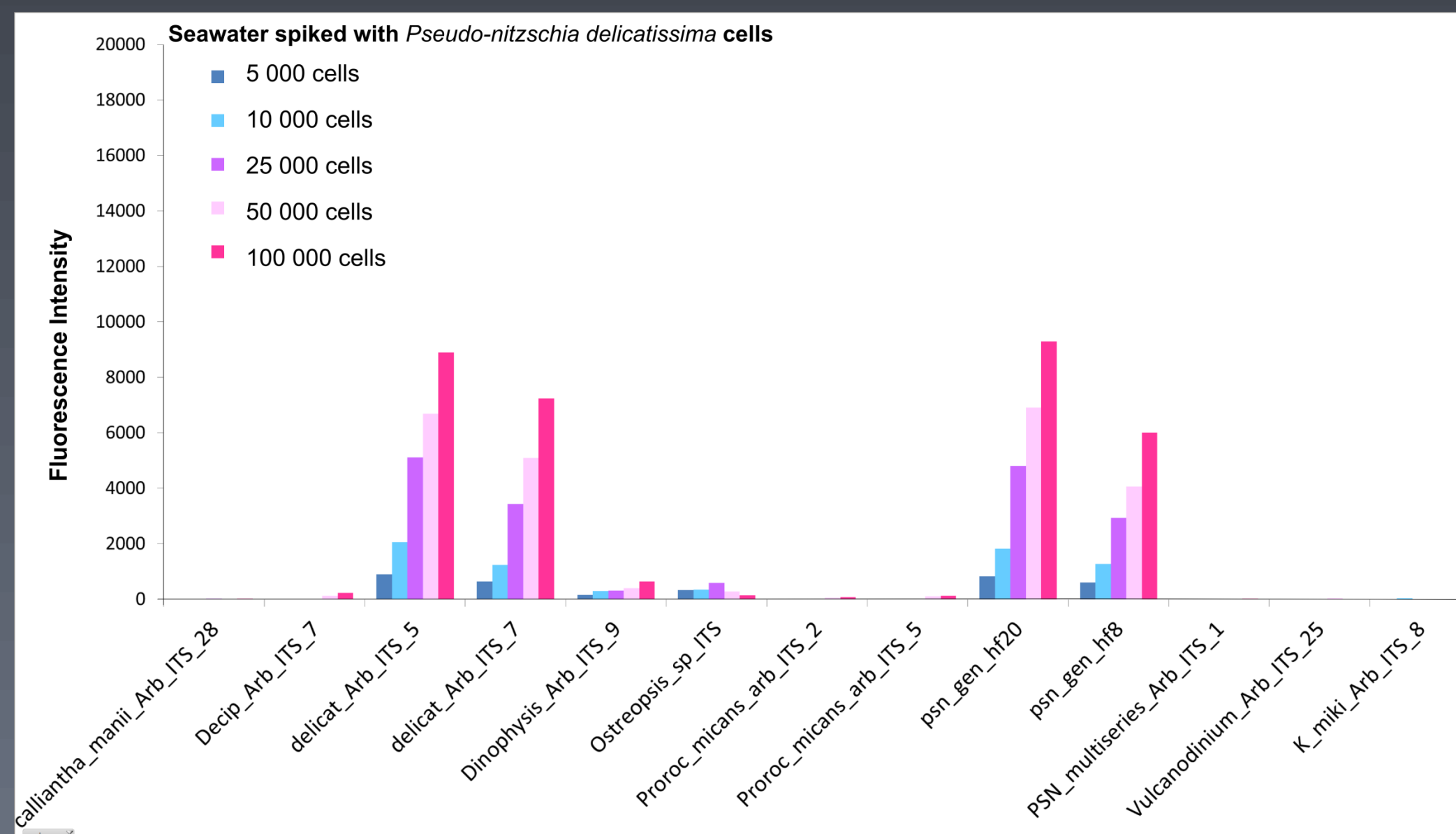
Detection of harmful algal blooms (HABs), also called red tides, has become a challenging concern due to the direct impacts on public health and economy. Current methods consist in microscopic identification and enumeration of the cells of interests. However it is time-consuming, tedious and requires expert taxonomists. Advances in molecular biology allow the development of new tools. In this context, we aimed at developing a new DNA microarray: the phytochip. It should be a rapid and accurate method dedicated to the identification of the main toxic phytoplankton species in French waters.

Results

P. multiseri hybridization: example of a microarray scan and probes intensity



Hybridization of seawater spiked with an increasing number of *P. delicatissima* cells



Specificity of the probes was tested with various monoclonal cultures, mixtures of DNA from cultures, and natural seawater spiked with *P. delicatissima* cells. 8 species of the *Pseudo-nitzschia* genus can be detected and accurately identified by specific probes of the phytochip. We propose the following hierarchical probe-set determination key to identify *Pseudo-nitzschia* species in environmental samples.

After amplification with the Eukaryotes primers, *Pseudo-nitzschia* were detected in Cabourg (Bay of Seine) in samples collected the 04/07/12 and 06/08/12. *P. americana* was detected weeks 27 and 32, *P. pungens* only week 32 and *P. australis* was detected week 27.

Perspectives

Specific probes were designed for several *Pseudo-nitzschia* species responsible for the ASP (Amnesic Shellfish Poisoning), species inducing PSP (Paralytic shellfish Poisoning) e.g. *Alexandrium* sp. and DSP (Diarrheic Shellfish Poisoning) e.g. *Dinophysis* sp. syndromes. We are currently testing the specificity and sensitivity of the phytochip using DNA samples from our collection. A series of environmental samples collected in the Bay of Seine will be analyzed and results will be compared with microscopic cell counts. The phytochip seems to be a promising method for the identification of toxic phytoplankton and could be a useful tool to help environmental monitoring.

Ludwig W, et al. (2004) ARB: a software environment for sequence data. Nucl Acids Res 32:1363–1371

LeBerre V, et al. (2003) Dendrimeric coating of glass slides for sensitive DNA microarrays analysis. Nucleic Acids Research 31

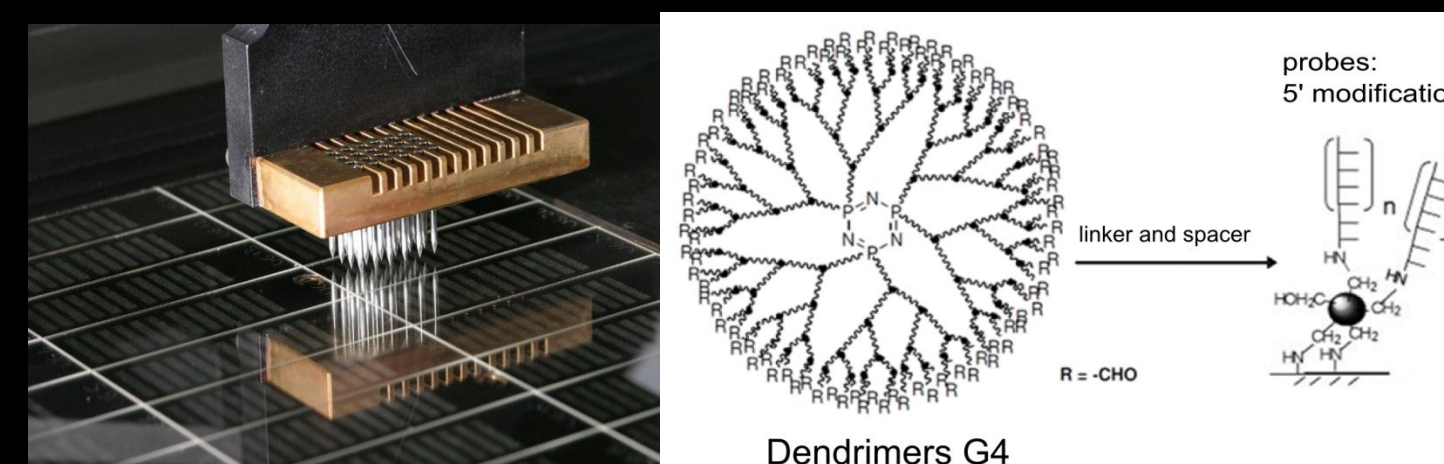
Trevisiol E, et al. (2003) Dendrimeric slides, dendrichips: a simple chemical functionalization of glass slides with phosphorus dendrimers as an effective means for the preparation of biochips. New Journal of Chemistry 27

Materials and Methods

1. Probe design : ARB software



2. Spotting on dendrimeric coating glass slides



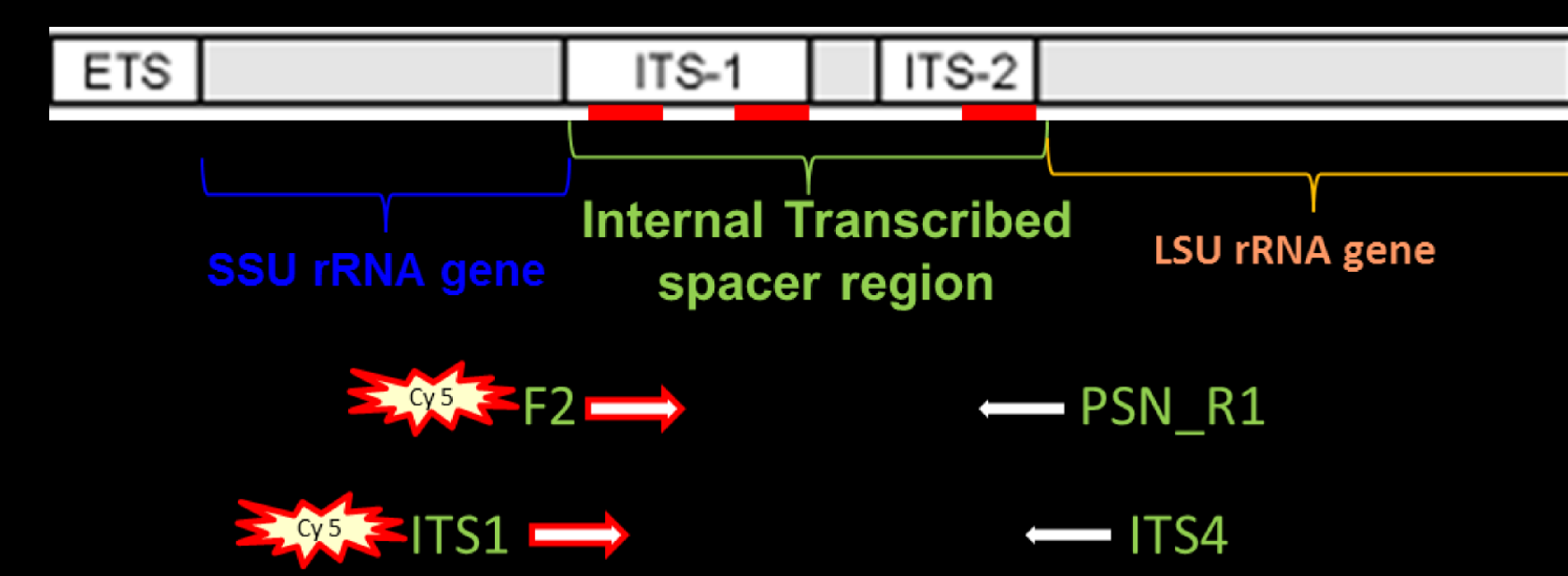
3. DNA extraction

(CTAB procedure or the DNAeasy Plant Mini Kit -Qiagen) :

- microalgae cultures
- seawater spiked with cells
- environmental samples

4. Asymmetric PCR with Cy5 labelled forward primer

(*Pseudo-nitzschia* genus and Eukaryotes)



5. Hybridization : 65° C for 30 min on 8 zone custom slides

6. Scan: Innoscan 900

7. Data analyses: MAPIX (Innopsys)

Probes were designed for the following species:

Pseudo-nitzschia

P. americana
P. australis
P. brasiliana
P. caciaantha
P. calliantha
P. decipiens
P. delicatissima
P. fraudulenta
P. galaxiae
P. Pseudodelicatissima / cuspidata
P. multiseri
P. pungens
P. seriata
P. subpacific



Alexandrium

A. catenella
A. minutum
A. tamarens
A. tamutum



Karenia

K. brevis
K. mikimitoi
K. selliformis



Dinophysis

D. acuta
D. acuminata
D. norvegica



and

Lingulodinium polyedrum
Ostreopsis ovata
Vulcanodinium rugosum



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