

Unveiling saxitoxins (STXs) synthesis potential of dinoflagellate Alexandrium through STXs synthesis genes (sxt) analysis

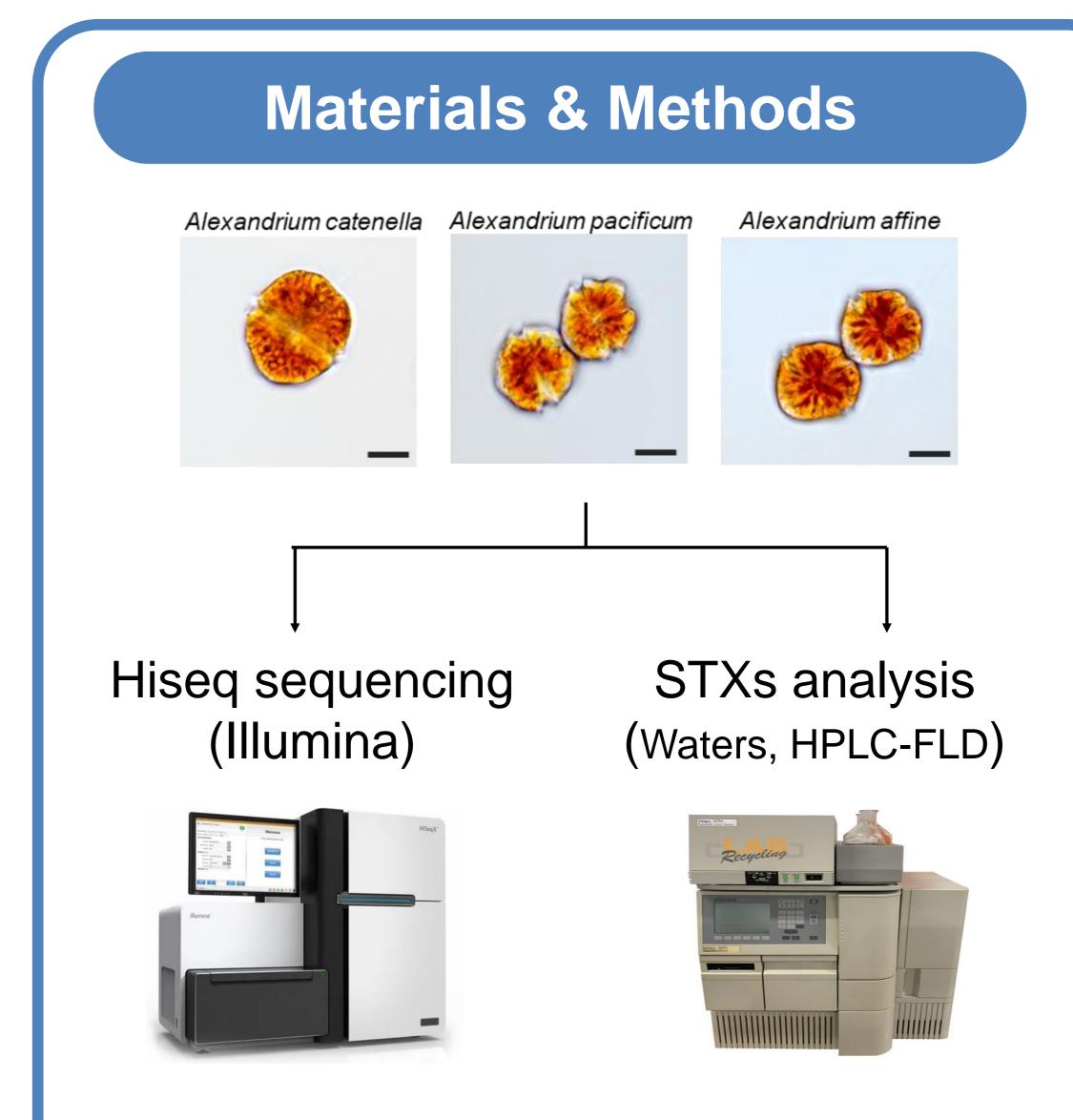
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Saxitoxins (STXs) producing dinoflagellates Sackground & Purpose Station Station (G') (B') Saxitoxins (STXs) Saxitoxins (STXs)

- Marine dinoflagellate *Alexandrium* occurs in oceans around the world, and some can form harmful algal blooms and produce toxic substances, <u>saxitoxins</u> (STXs).
- <u>STXs synthesis genes (sxt)</u> are known to be important for STXs production ability, however, it remains controversial.
- In the present study, <u>we identified several sxt genes</u> from toxic and nontoxic *Alexandrium* species through transcriptome analysis.
- Physiological and transcriptional responses of *A. affine* under various nutrient and temperature conditions were analyzed



 A. catenella, A. pacificum and A. affine were obtained from Korea Institute of Ocean Science & Technology (KIOST, Jangmok, Korea).

Results sxtG sxtA tamarense - UAT-014-009 60 A. affine - Alex02 A --- Akanthomyces lecanii 66 Pyrodinium bahamense Microseira wollei Heteroscytonema crispum Aphanizomenon flos-aquae - Corallococcus aberystwythensis Dolichospermum circinale - Catellatospora bangladeshensis Cylindrospermopsis raciborskii Raphidiopsis brookii Microcvstis aeruginosa or Leptolyngbya sp. 100 Anaeromyces robustus Cyanobacteria bacterium 98 - Acaryochloridaceae cyanobacterium Acaryochloris marina Dinoflagellate Vyanobacteria

Figure 1. A maximum likelihood (ML) tree of core enzymes sxtA4 and sxtG identified from *Alexandrium* comparing with diverse organisms. <u>Each enzymes are phylogenetically conserved among toxic dinoflagellates.</u>

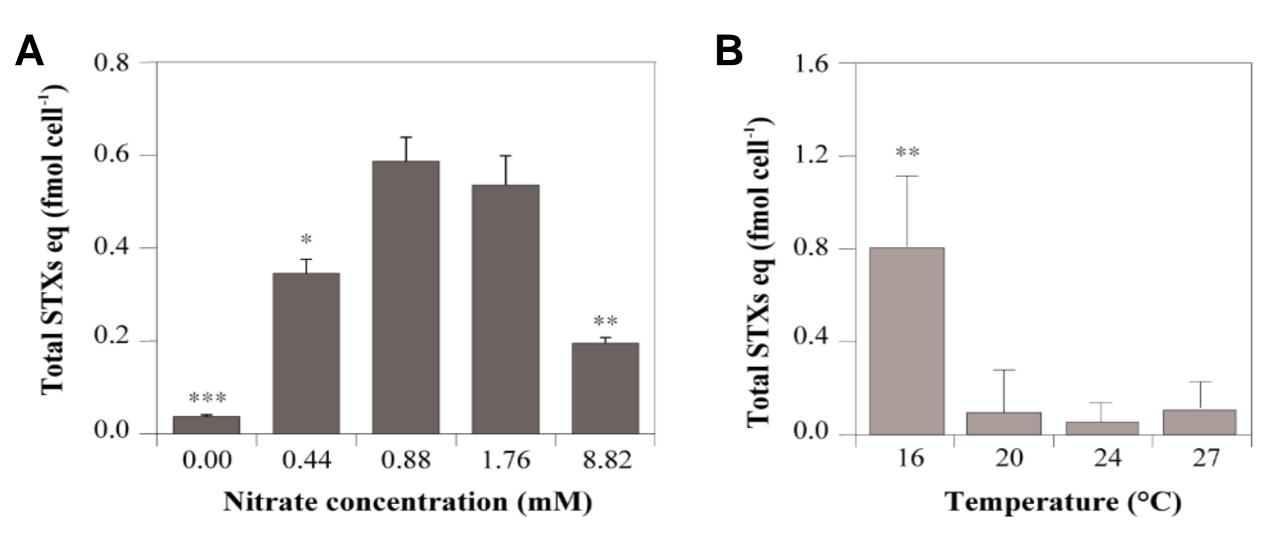


Figure 2. Total toxicity of *Alexandrium affine* cultured under various nitrate concentrations (A) and water temperatures (B). <u>STXs level of *A. affine* is up to 100-fold lower</u> than that of *A. pacificum* and *A. catenella*, and <u>toxin levels are varied by environmental changes.</u>

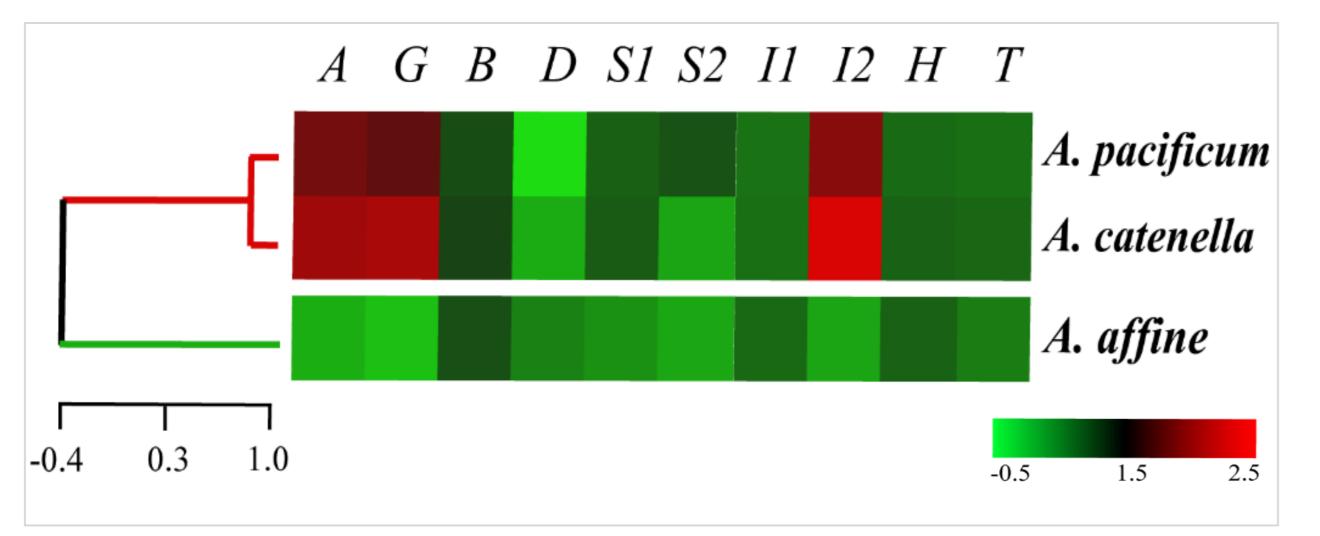
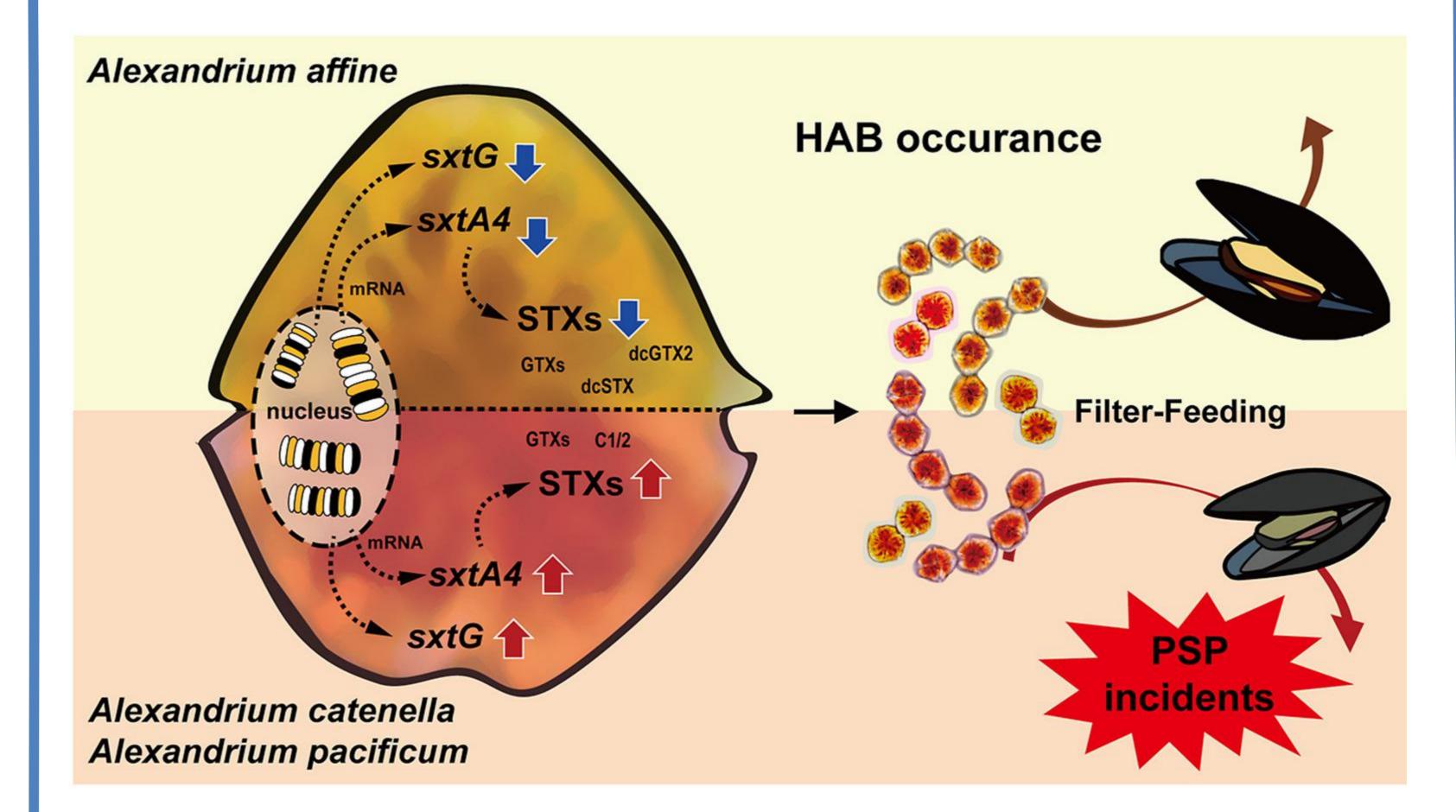


Figure 3. A heatmap of *sxt* RPKM (Reads Per Kilobase of transcript, per Million mapped reads). Expressional patterns of *sxt* were varied among toxic and non-toxic species, with <u>significantly lower expression levels of *sxtA*, *sxtG*, and *sxtl* in *A. affine*.</u>

Graphical Abstract & Conclusion



- A. affine is well known as a non-toxic species, with the exception of certain strains from Vietnam (Nguyen-Ngoc, 2004) and the Philippines.
- Similarly, STXs were detected in *A. affine* isolated from the southern coast of Korea, and its total STXs were significantly lower (<0.8 STXs eq fmol/cell) than those of toxic *A. catenella* and *A. pacificum*.
- STXs biosynthesis *sxt* genes were identified in *A. affine* through transcriptome analysis, transcript levels of the *sxtA*, *sxtG* and *sxtI* of *A. affine* were extremely low.
- These suggest that *A. affine* has the potential to produce STXs, however, the toxicity is much lower or negligible, making it less likely to cause PSP incidents in marine environments.





