

ANATOXIN-A IN SEVERAL FRESHWATER LAKES IN FRANCE: OCCURRENCE AND PHYLOGENY OF BIOSYNTHESIS GENES



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Cyanobacteria and cyanotoxins in France

Many freshwaters experiencing heavy

cyanobacterial blooms





The microcystins, the only cyanotoxins monitored by the Health Agency in France



The bloom-forming species are known to produce other cyanotoxins especially neurotoxins

What about anatoxin-a?



Dog poisonings in French rivers

(Gugger et al. 2005, Cadel-Six et al. 2009)

Many lakes with potentially anatoxin-a producing species (e.g Dolichospermum, Aphanizomenon)





ELSEVIER

Toxicon 45 (2005) 919-928

TOXICON

www.elsevier.com/locate/toxicon

First report in a river in France of the benthic cyanobacterium *Phormidium favosum* producing anatoxin-a associated

with dog neurotoxicosis

Muriel Gugger^{a,*}, Séverine Lenoir^{a,b}, Céline Berger^a, Aurélie Ledreux^a, Jean-Claude Druart^c, Jean-François Humbert^c, Catherine Guette^a, Cécile Bernard^a

Recent reports in freshwater lakes in Europe (*e.g* Rantala-Ylinen *et al.* 2011, Shams *et al.* 2015) but none in France



A three-year survey in Lake Aydat



- A natural lake aged of 8000 years
- Hypereutrophic state



Dolichospermum (aka Anabaena) macrospora
blooms every year in Autumn

Sampling in the water column in September-October



A three-year survey in Lake Aydat



Detection of anatoxin-a by LC-MS-MS analyses





Detection of anaC gene by PCR amplification

Anatoxin-a biosynthetic gene (*ana*) cluster of *Anabaena sp.* strain 37 (Rantala-Ylinen *et al.* 2011)





Sabart et al. Harmful algae 2015

Occurrence of ana genes at a larger scale



Occurrence of ana genes at a larger scale



Widespread occurrence of ana genes among the studied lakes



A phylogenetic study of anatoxin-a genes at a larger scale

High number of new sequences underlying the diversity of potentially producing strains

A phylogenetic study of anatoxin-a genes at a larger scale

Maximum likelyhood tree of anaC gene (98% identity)





anaC sequences from our study are clustering with Nostocales

A phylogenetic study of anatoxin-a genes at a larger scale

0.01 Anabaena-37-JF803645 -227-1-13 Anabaena-54 Apha-issatschenkoi-KM245025 409-3 Aphanizomenon-3 Cuspidothrix-issatschenkoi-LBRI48-KM245023 Nostocales Cuspidothrix-issatschenkoi-RM-6-KM245024 --- Uncultured-cyanobacterium-AB638256 -Cvlindrospermum-stagnale-PCC7417-CP003642 -Oscillatoria-PCC6506-clone3-AY768507 Oscillatoria-PCC6506-clone5-AY768508 227-2-1 -227-1-9 -227-2-11 _____227-2-4 227-1-11 227-2-13 -227-2-2 -227-1-10 -227-1-6 -Oscillatoria-193 —Phormidium-favosum-PMC240-04 hormidium-uncinatum-JX088084 439-16 141-aa <u>441-e</u> 439-е 439-s 437-7 -438-3 -437-23 437-26 437-1 one predominant _439-f 439-6 **Oscillatoriales** « ubiquitous » sequence -439-5 _____439_7

Maximum likelyhood tree of the anaF gene (100% identity)

anaF sequences from our study are clustering with Oscillatoriales

Conclusions



To improve our knowledge of anatoxin-a producers in freshwater lakes:









Isolation of anatoxin-a potentially producing strains

- Akinetes germination experiment
- Screening for the presence of *ana* genes
- Confirmation of anatoxin-a production by LC-MS-MS analyses

WORK IN PROGRESS...



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Thank you for your attention

