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Neurotoxic cyanobacterial toxins

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ABSTRACT

Worldwide development of cyanobacterial blooms has significantly increased in marine and continental waters in the last century due to water eutrophication. This phenomenon is favoured by the ability of planktonic cyanobacteria to synthesize gas vesicles that allow them to float in the water column. Besides, benthic cyanobacteria that proliferate at the bottom of lakes, rivers and costal waters form dense mats near the shore. Cyanobacterial massive proliferation is of public concern regarding the capacity of certain cyanobacterial strains to produce hepatotoxic and neurotoxic compounds that can affect public health, human activities and wild and stock animals. The cholinergic synapses and voltage-gated sodium channels constitute the targets of choice of cyanobacterial neurotoxins. Anatoxina and homoanatoxin-a are agonists of nicotinic acetylcholine receptors. Anatoxin-a(s) is an irreversible inhibitor of acetylcholinesterase. Saxitoxin, kalkitoxin and jamaicamide are blockers of voltage-gated sodium channels, whereas antillatoxin is an activator of such channels. Moreover the neurotoxic amino acid L-beta-N-methylamino-L-alanine was shown to be produced by diverse cyanobacterial taxa. Although controversial, increasing in vivo and in vitro evidence suggest a link between the ingestion of L-beta-N-methylamino-Lalanine and the development of amyotrophic lateral sclerosis/Parkinsonism-dementia complex, a neurodegenerative disease. This paper reviews the occurrence of cyanobacterial neurotoxins, their chemical properties, mode of action and biosynthetic pathways.

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1. Introduction

Cyanobacteria, the oldest known fossils (2.15 billions years) (Rasmussen et al., 2008), are photosynthetic ubiquitous micro-organisms that played a key role in the oxygenation of earth's atmosphere. Cyanobacteria constitute the most important nitrogen fixing organisms and possess the ability to harvest solar irradiance at wavelengths that most of photosynthetic organisms are unable. Cyanobacteria can occupy any terrestrial ecosystem including freshwater, marine and soil environments, as well as extreme habitats such as desserts, hot spring waters, and Artic and Antarctic environments (Sompong et al., 2005; Taton et al., 2006). Cyanobacterial massive proliferation, when dominated by toxic cyanobacteria, is of serious concern for public health, human activities, and animal life. Indeed, certain species of cyanobacteria are able to synthesize hepatotoxins, cytotoxins and neurotoxins (Carmichael, 1994; Sivonen and Jones, 1999; Briand et al., 2003). The lethal effects of cyanobacterial blooms upon stock animals were first described by Francis (1878) in a lake of the estuary of the Murray (Australia).

Cyanobacteria produce important hepatotoxins e.g., microcystins, nodularins and cylindrospermopsins. Microcystins are cyclic heptapeptides that possess the





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general structure (-p-Ala-X-p-MeAsp-Z-Adda-p-Glu-Mdha-). where X and Z are variable L-amino acids. D-MeAsp is D-erythro-β-methyl aspartic acid, Mdha is *N*-methyldehydroalanine, and Adda is 3-amino-9-methoxy-2,6,8-trimethyl-10-phenyldeca-4,6-dienoic acid (Sivonen and Iones, 1999). There exists more than 80 identified microcystin variants with different degrees of toxicity (i.p. mouse LD_{50} ranging from 50 to >1200 µg kg⁻¹) (Hotto et al., 2007). Microcystins are almost exclusively produced by planktonic cyanobacterial members of the genera Microcystis, Anabaena, Planktothrix, and in a small proportion by members of the genera Anabaenopsis, Hapalosiphon and Nostoc (Sivonen and Jones, 1999; Oksanen et al., 2004). Microcystins are powerful liver tumor promoters and potent inhibitors of the serine/threonine protein phosphatase-1 and -2A (Sivonen and Jones, 1999; Maynes et al., 2006). Nodularin is a cyclic pentapeptide hepatotoxin that differs from microcystins by lacking the amino acids D-Ala and X and by having L-Arg and N-methyldehydrobutyrine (Mdhb) instead of the variable amino acid Y and Mdha, respectively (Rinehart et al., 1988). Nodularin is produced by the bloom forming strain Nodularia spumigena, a hepatotoxic cyanobacterium that occurs worldwide in brackish waters, and by Nodularia harveyana PCC 7804 (Sivonen and Jones, 1999; Koskenniemi et al., 2007; Saito et al., 2001). Nodularin is a potent tumor promoter and inhibits serine/ threonine protein phosphatase-1 and -2A (i.p. mouse LD₅₀: 50–70 μ g kg⁻¹) (Gulledge et al., 2003). Cylindrospermopsin is a cytotoxic alkaloid consisting of a tricyclic guanidine moiety combined with hydroxymethyluracil that inhibits the synthesis of protein and of glutathione leading to cell death (i.p. CH_3 mouse LD_{50} : 2.1 mg kg⁻¹) (Ohtani et al., 1992; Runnegar et al., 2002). Cylindrospermopsin is produced by Cylindrospermopsis raciborskii, Aphanizomenon ovalisporum, Aphanizomenon flos-aquae, Umezakia natans, Raphidiopsis curvata and Anabaena bergii (Preussel et al., 2006). Cyanobacterial hepatotoxins are a serious threat for domestic and wild animal life as well as for human health (Carmichael et al., 2001; Koskenniemi et al., 2007; Hawkins et al., 1985). The World Health Organization has established a provisional drinking-water guideline for microcystin-LR: 1 mg l⁻¹ (World Health Organization, 2004).

Cyanobacterial neurotoxins target (i) cholinergic synapses: anatoxin-a and homoanatoxin-a are potent agonists of muscular and neuronal nicotinic acetylcholine receptor (nAChR) subtypes (Spivak et al., 1980; Thomas et al., 1993; Wonnacott et al., 1992), while anatoxin-a(s) is a potent irreversible inhibitor of acetylcholinesterase (Mahmood and Carmichael, 1986b) and (ii) sodium channels: saxitoxins are a group of structurally related molecules that block voltage-gated sodium channels (for review, see Llewellyn, 2006). The lipopeptides purified from marine cyanobacteria, kalkitoxin and jamaicamides A, B and C, also block voltage-gated sodium channels (LePage et al., 2005; Edwards et al., 2004), while antillatoxins A and B activate them (Li et al., 2001). Additionally, the nonprotein neurotoxic amino acid L-beta-N-methylamino-Lalanine, produced by diverse taxa of cyanobacteria, may be associated with the development of the amyotrophic lateral sclerosis/Parkinsonism-dementia complex, a neurodegenerative disease (Cox et al., 2003). The purpose of this paper

is to review the occurrence of cyanobacterial neurotoxins, their chemical properties, mode of action and biosynthetic pathways.

2. Neurotoxic alkaloids in freshwater cyanobacteria

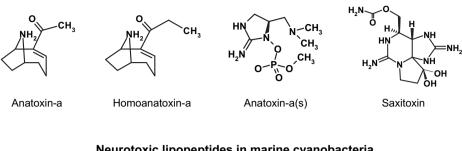
2.1. Anatoxin-a and homoanatoxin-a

(2-acetyl-9-azabicyanatoxin-a The neurotoxins clo[4.2.1]non-2-ene; UV λ_{max} at 227 nm; $\varepsilon = 12,000$; $C_{10}H_{15}NO$; EI-MS m/z [M]⁺ 165; Fig. 1) and its methylene homologue homoanatoxin-a (2-(propan-1-oxo-1-yl)-9-azabicyclo[4.2.1]non-2-ene; UV λ_{max} at 230 nm; C₁₁H₁₈NO; EI-MS m/z [M]⁺ 179; Fig. 1), are low molecular weight bicyclic secondary amines exclusively produced by cyanobacteria (Devlin et al., 1977; Skulberg et al., 1992). Anatoxin-a is synthesized by various members of the genera Anabaena (Harada et al., 1989), Aphanizomenon (Selwood et al., 2007), Cvlindrospermum (Sivonen et al., 1989), Microcvstis (Park et al., 1993), Oscillatoria (Sivonen et al., 1989), Planktothrix (Viaggiu et al., 2004) and Raphidiopsis (Namikoshi et al., 2003). Homoanatoxin-a is synthesized by some species corresponding to the genera Oscillatoria (Skulberg et al., 1992), Anabaena (Furey et al., 2003a), Raphidiopsis (Namikoshi et al., 2003) and Phormidium (Wood et al., 2007). Simultaneous synthesis of anatoxin-a and homoanatoxina was demonstrated for Raphidiopsis mediterranea Skuja (Namikoshi et al., 2003), and the axenic Oscillatoria PCC 9029 (Aráoz et al., 2005). The presence of anatoxin-a and homoanatoxin-a in several axenic cyanobacterial species exempt of bacterial contaminants - of the genera Anabaena (Rouhiainen et al., 1995) and Oscillatoria (Aráoz et al., 2005; Cadel-Six et al., 2007) confirmed the cyanobacterial origin of these toxins.

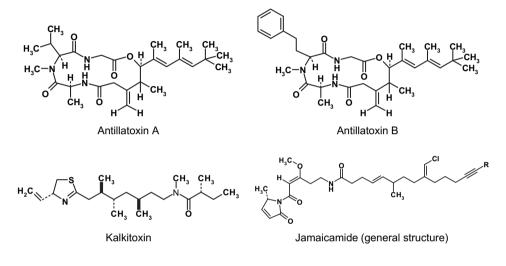
Massive proliferation of benthic neurotoxic cyanobacteria producing anatoxin-a and/or homoanatoxin-a near the shore of lakes and rivers has been shown to be fatal for wild and domestic animals, e.g., cows in Canada (Carmichael and Gorham, 1978), dogs in Scotland (Edwards et al., 1992), France (Gugger et al., 2005; Cadel-Six et al., 2007) and New Zealand (Wood et al., 2007), and Lesser flamingos in Kenya (Krienitz et al., 2003).

Electrophysiological experiments showed that anatoxin-a is a potent agonist of the muscle-type $\alpha 1_2\beta\gamma\delta$ nAChR. Thus, anatoxin-a induced neuromuscular blockade (of the depolanzing type), contracture of the frog's rectus abdominis muscle, depolarization of the frog sartorius muscle, desensitization, and alteration of the action potential (Spivak et al., 1980). Later, Thomas et al. (1993) working with chicken $\alpha 4\beta 2$ nAChR subunits expressed on mouse M 10 cells and chicken a7 nAChR expressed in oocytes from Xenopus laevis, showed that anatoxin-a is also a potent agonist of neuronal nAChR. Anatoxin-a, formerly known as "Very Fast Death Factor", kills mice 2-5 min after intraperitoneal injection preceded by twitching, muscle spasm, paralysis and respiratory arrest (i.p. mouse LD₅₀: 250 μg kg⁻¹; Devlin et al., 1977). Anatoxin-a is a depolarizing neuromuscular blocking agent. Neuromuscular blockade induced by anatoxin-a results from muscle membrane depolarization and desensitization. The binding of anatoxin-a to the nAChR induces the opening of the

Neurotoxic alkaloids in freshwater cyanobacteria



Neurotoxic lipopeptides in marine cyanobacteria



Cyanobacterial neurotoxic amino acid



L-β-*N*-methylamino-L-alanine (L-BMAA))

Fig. 1. Structure of the cyanobacterial neurotoxins.

receptor's channel allowing positively charged ions to move across it. This results in membrane depolarization. Prolonged exposure to anatoxin-a causes the desensitization of the nAChR that ultimately leads to the blockade of neuromuscular transmission (Spivak et al., 1980). Homoanatoxin-a was shown to be a potent analogue of anatoxina (Wonnacott et al., 1992). Additional mechanisms of toxicity were observed by systemic perfusion of sub-lethal and lethal doses of anatoxin-a to mice. Anatoxin-a seriously impaired blood pressure, heart rate and gas exchange (pO_2) and pCO₂) causing hypoxia and respiratory arrest, and severe acidosis that accompanied animal death (Adeymo and Sirén, 1992). Also, anatoxin-a induces the contraction of frog rectus abdominis muscle, as well as guinea pig ileum. In the latter case, the activation of nAChR from ganglionic interneurons by anatoxin-a, induced the release of acetylcholine that in cascade stimulated the muscarinic acetylcholine receptors leading to ileal contractions (Gordon et al., 1992).

The novel 9-azabicyclo[4.2.1]nonane semi rigid skeleton, and the potential pharmacological applications of anatoxin-a have led numerous organic chemists to develop diverse strategies for total synthesis of anatoxin-a (for reviews see Mansell, 1996; Parsons et al., 2000; Brenneman et al., 2004; Roe and Stockman, 2008). A series of eighteen anatoxin-a analogues were synthesized to use them as probes for characterizing the binding sites of anatoxin-a in muscle and neuronal types of nAChRs (Swanson et al., 1991; Wonnacott et al., 1991). These studies confirmed the important role that the side-chain stereochemistry of anatoxin-a plays in the binding of this toxin to the nAChR. Thus, the series of anatoxin-a analogues showed varying degrees of specificity for neuronal acetylcholine receptor subtypes. Actually, the anatoxin-a side-chain chemistry led to the synthesis of the methylene homologue named homoanatoxin-a (Wonnacott et al., 1991), before being purified from an oscillatorian strain (Skulberg et al., 1992).

Mass spectrometry coupled to gas or liquid chromatography is currently used for anatoxin-a and homoanatoxin-a detection at nanogram and picogram levels from water samples and cyanobacterial extracts. Liquid chromatography-tandem mass spectrometry and nanoelectrospray hybrid guadrupole time-of-flight mass spectrometry provide, reliability, sensitivity, selectivity, and structural and quantitative information for the analysis of anatoxin-a and homoanatoxin-a from cultured cyanobacteria and environmental samples (Himberg, 1989; Harada et al., 1993; Furey et al., 2003b; Maizels and Budde, 2004: James et al., 2005). Matrix-assisted laser desorption/ ionization time-of-flight mass spectrometry has also been applied to the detection of the low molecular mass anatoxin-a (165 Da) and homoanatoxin-a (179 Da) directly on cyanobacterial filaments of axenic strains of the genus Oscillatoria (Aráoz et al., 2008).

Feeding experiments with Anabaena flos-aquae and Oscillatoria formosa (producers of anatoxin-a and homoanatoxin-a, respectively) using¹³C-labeled precursors, showed that both, anatoxin-a and homoanatoxina biosynthesis proceed from acetate and glutamate (Hemscheidt et al., 1995). The source of carbons atoms is the same for anatoxin-a and homoanatoxin-a except for C-12. These results allowed Hemscheidt et al. (1995) to hypothesize the existence of a common precursor for both toxins which, following decarboxylation, may give rise to: i) anatoxin-a through enzymatic reduction of the common precursor, or to ii) homoanatoxin through enzymatic addition of a C₁ unit to the common precursor from the methyl donor S-adenosyl-L-methionine. Later, Namikoshi et al. (2004) identified L-methionine as the biosynthetic precursor of the methyl group at C-12 of homoanatoxina using L-[methyl-¹³C]-methionine in the culture of thecyanobacterium *Raphidiopsis mediterranea* (Fig. 2). Recently, Selwood et al. (2007) identified and chemically characterized for the first time the 11-carboxyl anatoxina derivative by high-resolution mass spectrometry in *Aphanizomenon issatschenkoi* (CAWBG02).

Note of the authors: as the present paper was under revision, a polyketide synthase coding sequence that is specific for strains producing anatoxin-a, or homoanatoxin-a was identified in *Oscillatoria* strain PCC 6506 (Cadel-Six et al., 2009). Partial genome sequencing of this cyanobacterium strain resulted in the identification of the putative gene cluster responsible for anatoxin-a and homoanatoxin-a production (Méjean et al., 2009). After gene annotation and deduction of their putative functions, Méjean et al. (2009) proposed a biosynthetic route for anatoxin-a and homoanatoxin-a. Moreover, feeding experiments using L-[U-¹³C-¹⁵N]Glu, L-[U-¹³C-¹⁵N]Arg, L-[U-¹³C-¹⁵N]Pro or L-[U-¹³C]Glu precursors, allowed the same authors to propose l-proline as the starter unit for anatoxin-a and homoanatoxin-a biosynthesis.

However, proline could be readily produced from either glutamate or arginine in the primary metabolism of cyanobacteria (Méjean et al., 2009). Therefore, we believe that more work needs to be done to gain direct insight into these neurotoxins biosynthetic route.

2.2. Anatoxin-a(s)

Anatoxin-a(s) (UV λ_{max} at 220 nm; C₇H₁₇N₄O₄P; HRFABMS [M + H]⁺ m/z 253.1066; Fig. 1) is an unusual natural organophosphate, structurally unrelated to anatoxin-a, that irreversibly inhibits acetylcholinesterase (Mahmood and Carmichael, 1986b). Its activity is similar to organophosphorous and carbamate insecticides such as paraoxon, physostigmine and pyridostigmine (Cook et al., 1988), and the chemical warfare agent sarin (Pita et al., 2003). Anatoxin-a(s) is produced by *Anabaena flos-aquae* strain NRC 525-17 (Mahmood et al., 1988) and *Anabaena lemmermannii* (Henriksen et al., 1997). Additionally,

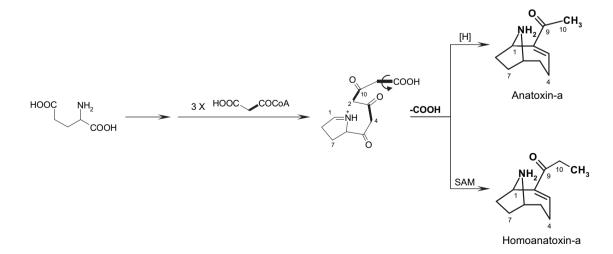


Fig. 2. Proposed biosynthetic route for homoanatoxin-a and anatoxin-a. Adapted, with permission, from Namikoshi et al. (2004). Copyright 2004 American Chemical Society. SAM: S-adenosyl-L-methionine.

acetylcholinesterase inhibition activity was found in extracts of *Anabaena spirolides*, isolated from environmental blooms of a water reservoir in Tapacurá, Brazil (Molica et al., 2005). When extracts of *A. flos-aquae* strain NRC 525-17 or purified anatoxin-a(s) were administered i.p. to mice, animal death was preceded by salivation, lacrimation, fasciculation, urinary incontinence and respiratory failure in 240 min (i.p. mouse LD_{50} : 20 µg kg⁻¹) (Mahmood and Carmichael, 1986b; Onodera et al., 1997a). Anatoxin-a(s) was involved in the poisoning of dogs, birds and swine in Canada (Mahmood et al., 1988), and wild birds in Denmark (Henriksen et al., 1997).

The Ellman test is currently used for the detection of anatoxin-a(s) (Mahmood and Carmichael, 1986b; Henriksen et al., 1997; Molica et al., 2005). Bachmann et al. (2000) adapted an electrochemical biosensor for the detection of anatoxin-a(s). The sensor system was composed of an Ag/AgCl reference electrode and a graphite working electrode in which acetylcholinesterase from *Drosophila* was immobilized. Acetylcholinesterase activity is monitored electrochemically from the formation of thiocholine following hydrolysis of acetylthiocholine chloride by the immobilized acetylcholinesterase. Devic et al. (2002) used a four-mutant set of *Drosophila* acetylcholinesterase to improve the specificity and the sensibility of the biosensor for anatoxin-a(s) detection.

Matsunaga et al. (1989) performed feeding experiments with A. flos-aquae NRC 525-17 using 50% ¹³C and 90% ¹⁵N to determine the chemical structure of this unique organophosphate. NMR analysis of the radioactive purified compound indicated that the N-CH₃ group was connected to the side chain of the imidazoline residue, and that the methylphosphate group was attached to one of the nitrogens through an ester bond. Anatoxin-a(s) slowly undergoes autolysis to give a mixture of a degradation product containing an imidazoline ring (High-Resolution Fast Atom Bombardment mass Spectrometry (HRFABS) $[M + H]^+$ m/z 159.1245) and monomethylphosphate that were separated by HPLC. The degradation product m/z159.1245 was converted into 4S-2-imino-4(dimethylaminomethyl)-imidazolidine (HRFABS $[M + H]^+$ *m*/*z* 143.1298) by catalytic hydrogenation. NMR analysis of both degradation products confirmed the stereochemistry of anatoxin-a(s) and the position of methylphosphate group at N₁. For detailed RMN data see Matsunaga et al. (1989) and Onodera et al. (1997a). The degradation compound 4S-2-imino-4(dimethylaminomethyl)-imidazolidine was synthesized starting from D- and L-asparagine in four steps by Matsunaga et al. (1989). However, total synthesis of anatoxin-a(s) has not yet been reported. Shimizu (1996) proposed a biosynthetic route for this natural organophosphate compound: the biosynthesis of anatoxin-a(s) may proceed from the unusual catabolism of arginine by a retro-Claisen condensation mechanism which involves the loss of a glycine moiety from an arginine molecule (Fig. 3).

2.3. Saxitoxin

Saxitoxin ($C_{10}H_{17}N_7O_4$; $M_r = 299.286480$; Fig. 1), mostly related with marine dinoflagellates outbreaks, was first purified from the Alaskan butter clam, *Saxidomus giganteus* in 1957 (Schantz et al., 1957). Shellfish accumulate saxitoxins through filter-feeding on toxic dinoflagellates. Saxitoxins are produced by some species of the genera *Alexandrium* (*A. andersoni, A. catenella, A. excavatum, A. fundyense, A. minutum, A. ostenfeldii, A. tamarense* and *A. tamiyavanichi*), *Gymnodinium catenatum* and *Pyrodinium bahamense* (see Llewellyn, 2006 for references). Some members of the freshwater cyanobacteria *Anabaena, Aphanizomenon, Cylindrospermopsis, Lyngbya* and *Planktothrix* were also reported to produce saxitoxins (see Table 1 for references). Freshwater mussels were also seen to accumulate saxitoxin when fed with toxic cyanobacteria (Negri and Jones, 1995).

Saxitoxin structure was determined by X-ray crystallography twenty years after its discovery (Schantz et al., 1975; Bordner et al., 1975). Soon later, Shimizu et al. (1976) isolated several new paralytic shellfish toxins, named gonyautoxins from shellfish samples and from the causative organism *Gonyaulax tamarensis*. Saxitoxins comprise a series of 30 structurally related natural molecules with two guanidinium moieties often referred to as paralytic shellfish poisoning. Structurally, saxitoxins are divided into two categories: saxitoxin and neosaxitoxin series. In each series, the structural variations are created by the presence and differential stereochemistry of the sulphate groups at C-11, and the occurrence of *N*-sulphate carbamoyl groups at C-17 (Shimizu, 1986; Negri et al., 1995, Fig. 4; Table 1).

Saxitoxins block sodium permeability of excitable membranes by reversible binding to specific amino acid residues located in the outer pore loops of the voltage-dependent sodium channels at site 1, as tetrodoxin does (Cestèle and Catterall, 2000; Bricelj et al., 2005). The blockage of voltage-gated sodium channels prevents the generation of a proper action potential in nerves and muscle fibres leading to neuromuscular paralysis and death by respiratory arrest. Saxitoxins are among the most toxic compounds known (i.p. mouse LD_{50} : 5–10 µg kg⁻¹; Schantz et al., 1975). The rank order of sodium channel blockade is

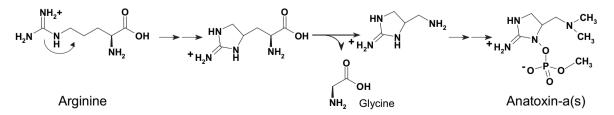


Fig. 3. Proposed anatoxin-a(s) biosynthetic pathway. Biosynthesis of anatoxin-a(s) involves the loss of glycine from arginine. Partially reprinted from Shimizu (1996), with permission from the *Annual Review of Microbiology*, Volume 50 © 1996 by Annual Reviews (www.annualreviews.org).

Table 1

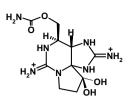
Saxitoxin analogues produced by freshwater cyanobacteria.

Toxin	Cyanobacterial strain	References
Saxitoxin (STX)	Anabaena circinalis	Negri et al., 1995; Teste et al., 2002
	Anabaena lemmermannii	Rapala et al., 2005
	Aphanizomenon flos-aquae	Ikawa et al., 1982; Mahmood and Carmichael, 1986a; Dias et al., 2002;
		Pereira et al., 2000
	Aphanizomenon gracile LMECYA40	Pereira et al., 2004
	Aphanizomenon issatschenkoi	Nogueira et al., 2004
	Cylindrospermopsis raciborskii	Lagos et al., 1999; Molica et al., 2002
	Cylindrospermopsis raciborskii C10	Castro et al., 2004
	Cylindrospermopsis raciborskii T3	Pomati et al., 2004
	Planktothrix sp.	Pomati et al., 2000
Decarbomoyl saxitoxin (dcSTX)	Anabaena circinalis	Negri et al., 1995
• · · ·	Aphanizomenon DC-1	Liu et al., 2006
	Aphanizomenon flos-aquae	Pereira et al., 2000; Dias et al., 2002
	Aphanizomenon issatschenkoi	Nogueira et al., 2004
	Cylindrospermopsis raciborskii C10	Castro et al., 2004
	Lyngbya wollei	Carmichael et al., 1997
Neosaxitoxin (neoSTX)	Aphanizomenon DC-1	Liu et al., 2006
	Aphanizomenon flos-aquae	Ikawa et al., 1982; Mahmood and Carmichael, 1986a; Dias et al., 2002;
		Pereira et al., 2000
	Aphanizomenon gracile LMECYA40	Pereira et al., 2004
	Aphanizomenon issatschenkoi	Nogueira et al., 2004
	Cylindrospermopsis raciborskii	Lagos et al., 1999; Molica et al., 2002
Gonyautoxin 1 (GTX1)	Aphanizomenon flos-aquae	Ferreira et al., 2001
Gonyautoxin 2 (GTX2)	Anabaena circinalis	Negri et al., 1995; Jones and Negri, 1997: Teste et al., 2002;
• • •	Cylindrospermopsis raciborskii	Lagos et al., 1999
	Cylindrospermopsis raciborskii C10	Castro et al., 2004
Gonyautoxin 3 (GTX3)	Anabaena circinalis	Negri et al., 1995; Jones and Negri, 1997; Teste et al., 2002
• • •	Aphanizomenon flos-aquae	Ferreira et al., 2001
	Cylindrospermopsis raciborskii	Lagos et al., 1999
	Cylindrospermopsis raciborskii C10	Castro et al., 2004
Gonyautoxin 4 (GTX4)	Aphanizomenon flos-aquae	Ferreira et al., 2001
Gonyautoxin 5 (GTX5)	Anabaena circinalis	Negri et al., 1995
	Aphanizomenon flos-aquae	Pereira et al., 2000; Dias et al., 2002
	Aphanizomenon issatschenkoi	Nogueira et al., 2004
Gonyautoxin 6 (GTX6)	Aphanizomenon flos-aquae	Pereira et al., 2000
	Cylindrospermopsis raciborskii	Molica et al., 2002
Decarbomoyl gonyautoxin 2 (dcGTX2)	Anabaena circinalis	Negri et al., 1995
	Lyngbya wollei	Carmichael et al., 1997; Onodera et al., 1997b
Decarbomoyl gonyautoxin 3 (dcGTX3)	Anabaena circinalis	Negri et al., 1995; Teste et al., 2002
	Aphanizomenon DC-1	Liu et al., 2006
	Lyngbya wollei	Carmichael et al., 1997; Onodera et al., 1997b
C1	Aphanizomenon flos-aquae	Ferreira et al., 2001
	Anabaena circinalis	Negri et al., 1995; Jones and Negri, 1997; Teste et al., 2002
	Cylindrospermopsis raciborskii T3	Pomati et al., 2004
C2	Anabaena circinalis	Negri et al., 1995; Jones and Negri, 1997; Ferreira et al., 2001; Teste et al., 2002
	Cylindrospermopsis raciborskii T3	Pomati et al., 2004
Lyngbya wollei toxin 1	Lyngbya wollei	Onodera et al., 1997b
Lyngbya wollei toxin 2	Lyngbya wollei	Onodera et al., 1997b
Lyngbya wollei toxin 3	Lyngbya wollei	Onodera et al., 1997b
Lyngbya wollei toxin 4	Lyngbya wollei	Onodera et al., 1997b
Lyngbya wollei toxin 5	Lyngbya wollei	Onodera et al., 1997b
Lyngbya wollei toxin 6	Lyngbya wollei	Onodera et al., 1997b

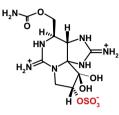
the following: carbamate saxitoxins > saxitoxin > neosaxi toxin > gonyautoxins > decarbamoyl saxitoxins > *N*-sulfocarbamoyl derivatives saxitoxins (Deeds et al., 2008; Indrasena and Gill, 1998).

A cyanobacterial bloom dominated by *Anabaena circinalis* in a dam near Forbes in central New South Wales (Australia) was suspected to cause the death of 14 sheeps. C-toxins, gonyautoxins and saxitoxin were found in extracts of *A. circinalis* and in the intestine contents of dead sheep (Negri et al., 1995). Saxitoxin producing cyanobacterial strains were also found in the water reservoir of Tapacurá (Brazil) supplying drinking water to 1.3 million inhabitants (Molica et al., 2005). Saxitoxin was

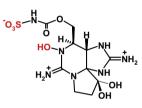
considered in the Schedule 1 of the Chemical Weapons Convention together with warfare agents such as mustard gas, sarin, ricin and many others (see review by Llewellyn, 2006 and references included). Several methods for saxitoxin detection have been developed which are highly sensitive and specific including fluorescence spectroscopy (Indrasena and Gill, 1998), HPLC with pre-column derivatization and fluorescence detection (Papageorgiou et al., 2005; Lawrence et al., 2005), radio-receptor binding assays (Ruberu et al., 2003), saxiphilin- (a saxitoxin binding protein) based methods (Llewellyn et al., 2001), immunodetection (Micheli et al., 2002) and cell-based assays (Humpage et al., 2007).



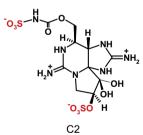
Saxitoxin (STX)



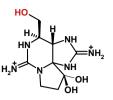
Gonyautoxin 2 (GTX2)



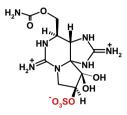
Gonyautoxin 6 (GTX6)



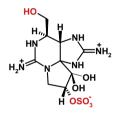
Lyngbya wollei toxin 4 (Lw toxin 4)



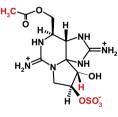
Decarbomoyl saxitoxin (dcSTX)



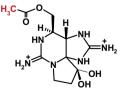
Gonyautoxin 3 (GTX3)



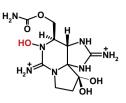
dcGonyautoxin 2 (dcGTX1)



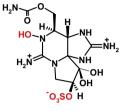
Lyngbya wollei toxin 1 (Lw toxin 1)



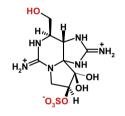
Lyngbya wollei toxin 5 (Lw toxin 5)



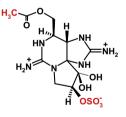
Neosaxitoxin (neoSTX)



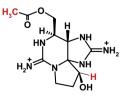




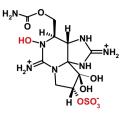
dcGonyautoxin 3 (dcGTX3)



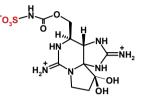
Lyngbya wollei toxin 2 (Lw toxin 2)



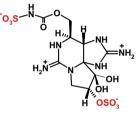
Lyngbya wollei toxin 6 (Lw toxin 6)



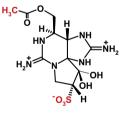
Gonyautoxin 1 (GTX1)



Gonyautoxin 5 (GTX5)



C1



Lyngbya wollei toxin 3 (Lw toxin 3)

Fig. 4. Saxitoxin analogues produced by some members of different cyanobacteria genera. From Llewellyn (2006). Reproduced by permission of The Royal Society of Chemistry (http://dx.doi.org/10.1039/b501296c). See Table 1 for species names and references.

Saxitoxin shows nanomolar affinity for certain voltagegated sodium channels types, and constitutes a suitable blueprint for developing new channel blockers for therapeutic purposes (Schantz and Johnson, 1992; Kohane et al., 2000; Garrido et al., 2007). Although the chemical synthesis of saxitoxin was achieved in the past (Tanino et al., 1977; Jacobi et al., 1984), novel synthetic pathways were designed to assemble the tricyclic skeleton that defines saxitoxin (Fleming and Du Bois, 2006; Fleming et al., 2007). Recently, the biosynthetic gene cluster of saxitoxin was discovered from the cyanobacterium *C. raciborskii* T3 (Kellmann et al., 2008). Saxitoxin gene cluster comprises more than 35 kb encoding for 31 open reading frames. By comparative sequence analysis, 30 enzymatic activities were assigned to 26 proteins of the saxitoxin synthase complex. The biosynthetic origin of saxitoxin comprises arginine, *S*-adeno-sylmethionine and acetate as it was proposed by Shimizu (1993) based on feeding experiments performed with *Alexandrium tamarense* and *Aphanizomenon flos-aquae*. The *in-silico* comparative DNA sequence analysis complemented

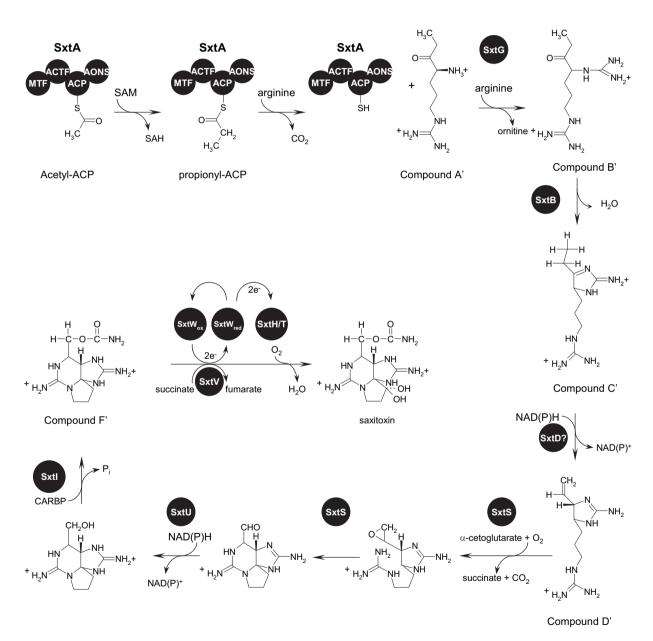


Fig. 5. Revised pathway for the biosynthesis of saxitoxin proposed by Neilan et al. From Kellmann et al. (*Appl. Environ. Microbiol.*, 2008, 74, 4044-4053 doi:10. 1128/AEM.00353-08) and reproduced with permission from American Society for Microbiology. The *sxt* genes from *Cylindrospermopsis raciborskii* T3 and their predicted functions: *sxtA*: methyltransferase; *sxtG*: amidinotransferase; *sxtB*: cytidine deaminase; *sxtD*: sterole desaturase-like protein; *sxtS*: phytanoyl-CoA dioxygenase; *sxtU*: alcohol dehydrogenase; *sxtI*: carbamoyltransferase; *sxtW*: ferredoxin; *sxtV*: succinate dehydrogenase; *sxtH*: phenylpropionate dioxygenase; *sxtT*: phenylpropionate dioxygenase.

with liquid chromatography–tandem mass spectrometry experiments carried out with extracts of *Anabaena circinalis* to detect the presence of the intermediate metabolites A', C' and E' allowed Neilan et al. to propose a revised pathway for saxitoxin biosynthesis (Fig. 5) (Kellmann et al., 2008).

3. Neurotoxic lipopeptides in marine cyanobacteria

3.1. Antillatoxin

Antillatoxin A ($C_{28}H_{45}N_3O_5$; UV λ_{max} at 230 nm; $\varepsilon = 12,000$; HRFABMS m/z [M + H]⁺ 504.3436; Fig. 1) is a structurally novel lipopeptide with a high degree of methylation produced by the tropical marine cyanobacterium Lyngbya majuscula collected in Curaçao (Orjala et al., 1995). The cyclic lipopeptide antillatoxin A is composed of a tripeptide that forms both ester and amide linkages with a highly methylated lipid section. For complete ¹H and ¹³C NMR data see Oriala et al. (1995). Yokokawa et al. (2000) and Li et al. (2004). In addition to antillatoxin A, specimens of L. majuscula collected from Collado Reef (Puerto Rico) and Bush Key, Dry Tortugas (Florida Keys, USA) were shown to produce an N-methyl homophenylalanine homologue called antillatoxin B (C₃₃H₄₈N₃O₅; UV λ_{max} at 209 nm, ε = 50,119 and at 240 nm, ε = 114,824.06; HRFABMS m/z $[M + Na]^+$ 566.3596; Fig. 1). For complete ¹H and ¹³C NMR data see Nogle et al. (2001).

The structure of antillatoxin A, initially formulated from spectroscopic information, was revised at one stereocenter (C-4) after its chemical synthesis by Yokokawa et al. (1999, 2000). The preferred stereochemistry of antillatoxin A was further confirmed as a result of synthesis of four different stereoisomers of antillatoxin A (all possible C-4 and C-5 diastereomers) (Li et al., 2004). Consequently, natural antillatoxin A has a (4R,5R)-configuration (Yokokawa et al., 2000; Li et al., 2004). Total synthesis of antillatoxin A and its stereoisomers was achieved by Yokokawa and Shioiri (1998), Yokokawa et al. (1999, 2000) and recently by Lee and Loh (2006).

Antillatoxin A was first characterized as one of the most ichthyotoxic compounds (Lethal Concentration 50 $(LC_{50}) = 0.1 \mu M$) exceeded only by brevetoxins (Orjala et al., 1995). Later, it was shown that antillatoxin A induced a rapid neuronal death in cerebellar granule cell cultures $(LC_{50} = 0.18 \mu M)$. The cytotoxicity of antillatoxin A was prevented by dextrorphan and MK-801, two non-competitive N-methyl-D-aspartic acid (NMDA) receptor antagonists (Berman et al., 1999). The work of Li et al. (2001) showed for the first time that the molecular target of antillatoxin A was the voltage-gated sodium channels. Indeed, they showed that neurotoxicity, as well as Ca^{2+} influx into cerebellar granule cells induced by antillatoxin A, was antagonized by tetrodotoxin, a well known sodium channel blocker. Furthermore, antillatoxin A was found to enhance Na⁺ influx in intact neurons, effect that was also antagonized by tetrodotoxin ($EC_{50} = 98.2 \text{ nM}$) confirming that antillatoxin A is an activator of voltage-gated sodium channels (Li et al., 2001).

Antillatoxin A is a member of the lipid-soluble gating modifier toxins of voltage-gated sodium channels that include brevetoxins, batrachotoxin, veratridine and gambierol (Cao et al., 2008). This group of toxins produces a rapid and concentration-dependent increase of intracellular Na⁺ in neocortical neurons that can be antagonized by tetrodoxin. Intracellular Na⁺ has been shown to modulate NMDA receptor activity. The increase of intracellular Na⁺ selectively up-regulates synaptic responses mediated by NMDA receptors but not by non-NMDA receptors (Yu and Salter, 1998; Yu, 2006). In the case of exposure of neocortical neurons to antillatoxin A (300 nM) the increase of intracellular Na⁺ exceeded 40 mM (Cao et al., 2008). Further work needs to be done in order to determine the binding site of antillatoxin A in voltage-gated sodium channels.

Antillatoxin B showed reduced sodium channel-activation properties ($EC_{50} = 1.77 \mu$ M) and exhibited less ichthyotoxic activity ($LC_{50} = 1 \mu$ M) compared to antillatoxin A, suggesting that the substitution of an *N*-methyl homophenylalanine residue for an *N*-methyl valine residue in antillatoxin B is responsible for its decreased activity (Nogle et al., 2001, Fig. 1). Similarly, the (*4R*,*5R*)-antillatoxin A is 25-fold more potent than its other three stereoisomers [(*4S*,*5R*)-, (*4S*,*5S*)- and (*4R*,*5S*)-antillatoxin A], indicating that the overall molecular topology of antillatoxin A is affected by changes at the stereocenter (C-4) (Li et al., 2004).

3.2. Kalkitoxin

Kalkitoxin (C₂₁H₃₈N₂OS; UV (MeOH) λ_{max} 250 nm; $\varepsilon = 2600$; High-resolution electron impact mass spectrometry m/z [M]⁺ 366.2696; Fig. 1), is a thiazoline-containing lipopeptide discovered and purified from organic extracts of Lyngbya majuscula collected in the coasts of Curaçao using bioassay guided fractionation (Wu et al., 2000). Later, Nogle and Gerwick (2003) showed that L. majuscula specimens collected in the shallow coasts of Puerto Rico also produced kalkitoxin. Natural kalkitoxin possesses a 2,4-disubstituted thiazoline, a lipophilic chain and an unsaturated CH₂=CH₂ unit. For complete ¹H NMR (benzene- d_6 , 500 MHz) and ^{13}C NMR (DMSO- d_6 , 100 MHz) see Wu et al. (2000). Based on NMR analysis, Wu et al. (2000) assigned four stereochemical possibilities to kalkitoxin. In order to determine the absolute stereochemistry of natural (3R, 7R, 8S, 10S, 2'R)-kalkitoxin, they synthesized all possible stereoisomers and compared their ¹³C NMR spectra. Total synthesis of (+)-kalkitoxin was also achieved by White et al. (2004).

Initial studies have shown that kalkitoxin was ichthyotoxic to the goldfish *Carassius auratus* ($LC_{50} = 700 \text{ nM}$) and toxic to the aquatic crustacean brine shrimp (*Artemia salina*) with an $LC_{50} = 170 \text{ nM}$ (Wu et al., 2000). Accordingly, cytotoxic studies performed with synthetic (+)-kalkitoxin and two synthetic precursors using the human colon cell line HCT-116 showed that the thiazoline moiety of kalkitoxin is required for cytotoxicity ($IC_{50} = 1.0 \times 10^{-3} \,\mu\text{g ml}^{-1}$) (White et al., 2004). Kalkitoxin was also shown to induce delayed neurotoxicity in cerebellar granule neurons in a concentration-dependent manner ($LC_{50} = 3.86 \text{ nM}$). The delayed kalkitoxin toxicity was prevented only when the NMDA receptor antagonists (dextrorphan and MK-801) were present during the 22 h post-exposure period (Berman et al., 1999). Kalkitoxin interaction with voltagegated sodium channels was demonstrated using cerebellar granule neurons in culture (LePage et al., 2005). Kalkitoxin blocks veratridine-induced intracellular elevation of Ca²⁺ and neurotoxicity in a concentration-dependent manner $(EC_{50} = 262.7 \text{ nM})$ showing indirectly that kalkitoxin blocks voltage-gated sodium channels. Furthermore, ligandbinding assay data using cerebellar granule cells showed that kalkitoxin alone did not interfere with [³H]-batrachotoxin binding to the sodium channels; however, in the presence of deltamethrin, a positive allosteric modulator of sodium channels, kalkitoxin inhibited [³H]-batrachotoxin binding to the voltage-gated sodium channels $(IC_{50} = 11.9 \text{ nM}; \text{ LePage et al., } 2005)$. Since batrachotoxin binds to voltage-gated sodium channels only when the channel is in its open conformation (Catterall et al., 1981), the latter results provide evidences that kalkitoxin acts as a blocking agent of voltage-gated sodium channels.

3.3. Jamaicamide

Jamaicamide A ($C_{27}H_{37}O_4N_2ClBr$; UV (MeOH) λ_{max} 272 nm; $\varepsilon = 7943$; HRFABMS $[M + H]^+$ at m/z 567.1625; Fig. 1) is a novel highly functionalized neurotoxic lipopeptide possessing an unusual alkynyl bromide, vinyl chloride, β -methoxy eneone system, and a pyrrolinone ring. Jamaicamide A and two other isomers, jamaicamide B $(C_{27}H_{37}O_4N_2Cl)$ and jamaicamide C $(C_{27}H_{39}O_4N_2Cl;)$ were purified and characterized from a dark green strain of the marine cyanobacterium Lyngbya majuscula (strain JHB) growing in low abundance in Hector's Bay, Jamaica. Jamaicamide B is a debromo analogue of jamaicamide A, while in jamaicamide C, which also lacks the bromine atom, a terminal olefin replaces the terminal alkyne of jamaicamide B. For ¹H and ¹³C NMR data of jamaicamides see Edwards et al. (2004). The exceptional structure of jamaicamide B that includes a vinyl chloride was exploited to visualize and distinguish L. majuscula strain "3L" (curacin A producer) from L. majuscula strain JHB (jamaicamides A, B and C producer) by MALDI-TOF imaging at m/z 374 for curacyn and at m/z 511 for jamaicamide B (Simmons et al., 2008).

The chemical structure of jamaicamides A, B and C suggested that these molecules derive from a mixture of polyketide (9 acetate units), amino acid (L-Ala, β -Ala) and methionine-derived methyl groups. Based on feeding experiments that strongly supported a mixed polyketide synthase/nonribosomal peptide synthetase assembly for jamaicamides, a 58 kb gene cluster of 17 open reading frames was cloned and the biosynthetic pathway of jamaicamides deciphered. The *jam* gene cluster from *L. majuscula* JHB is organized in a remarkably co-linear arrangement with respect to its proposed biosynthesis (Fig. 6) (Edwards et al., 2004).

The jamaicamides A, B and C showed similar cytotoxicity to H-460 human lung and Neuro-2a mouse neuroblastoma cell lines ($LD_{50} = 15 \mu M$). Jamaicamides were tested for their capacity either to activate or block voltage-gated sodium channels using an antagonism cell bioassay (Manger et al., 1995) with cerebellar granule neurons, ouabaine, veratridine, brevetoxin and saxitoxin. All jamaicamide isomers showed channel-blocking activity at 5 μ M. None of the jamaicamide isomers exhibited sodium channel-activating activity (Edwards et al., 2004).

4. Neurotoxic amino acid

4.1. L-Beta-N-methylamino-L-alanine

The non-protein amino acid L-beta-N-methylamino-L-alanine (L-BMAA; 2-amino-3-methylaminopropanoic acid; C₄H₁₀N₂O₂; MW: 118.1344; Fig. 1) was initially isolated from highly toxic seeds of the false sago palm, the gymnosperm Cycas circinalis L. (Vega and Bell, 1967; Vega et al., 1967). A possible link was established between I-BMAA and the high incidence of a progressive neurological disease displaying clinical symptoms and histological aspects similar to amyotrophic lateral sclerosis/Parkinsonism-dementia complex (ALS/PDC), among the Chamorro people who traditionally used cycad seeds as a source of food and medicine in the Mariana islands of Guam and Rota (Spencer et al., 1986; Spencer et al., 1987a). Previous studies showed a correlation between exposure to cycad seeds and motor neurone disease in West New Guinea in Indonesia (Spencer et al., 1987b), and in Kii peninsula of Honshu island in Japan (Spencer et al., 1987c). Neurological deficits observed following repeated oral administration of L-BMAA to macaques (Macaca fascicularis) supported the hypothesis that cycad seeds might play a significant role in the aetiology of ALS/PDC in Guam and elsewhere (Spencer et al., 1986, 1987a). However, other experiments with primates fed on cycad flour (with unknown concentrations of L-BMAA) did not produce any clinical signs of neurological diseases (Duncan et al., 1988; Garruto et al., 1988, 1991).

Recently, the possibility that L-BMAA could trigger neurodegenerative disease has been put again at the forefront following the publications by Cox et al. who proposed biomagnification of cyanobacterial L-BMAA as a mechanism to generate large amount of neurotoxins in the Guam ecosystem (Cox and Sacks, 2002). This hypothesis was corroborated by (i) the discovery that L-BMAA occurs as both, free and protein-bound forms throughout tissues of flying foxes (Pteropus mariannus mariannus). Flying foxes feed on cycads (Cycas micronesica) and constitute a component of the traditional Chamorro diet (Banack et al., 2006), and (ii) the demonstration that diverse taxa of cyanobacteria including Nostoc symbionts isolated from coralloid roots of C. microsenica synthesize L-BMAA (Cox et al., 2003, 2005). Moreover, protein-bound L-BMAA was found to remain in Chamorro foods prepared from cycad flour (Banack et al., 2006). In Peru, people seasonally collect in highland lakes globular colonies of Nostoc commune, called "llullucha". Such colonies purchased in different markets from Cuzco (South East of Peru) were analyzed by UPLC-MS and LC/MS/MS and were found to contain L-BMAA (Johnson et al., 2008). Recent reports also described the presence of L-BMAA in a marine cyanobacterium isolated from the island of Oahu in Hawaii (Banack et al., 2007) and in cyanobacterial cultures representing the taxonomic diversity and geographic distribution in Southern Africa (Esterhuizen and Downing, 2008). Co-occurrence of

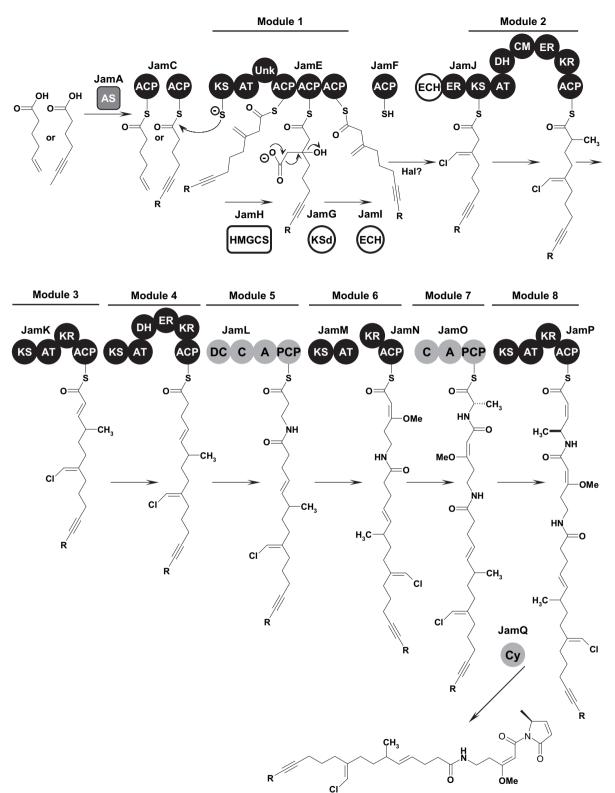


Fig. 6. Jamaicamide biosynthetic pathway. Reproduced, with permission, from Edwards et al. (2006). Copyright © 2004 Elsevier Ltd. The Jam proteins from *Lyngbya majuscula* JHB *jam* gene cluster and their predicted functions: JamA: hexanoyl-ACP synthetase; JamC: acyl carrier protein; JamE: polyketide chain extension; JamF: acyl carrier protein; JamG: ketosynthase/decarboxylase; JamH: 3-hydroxy-3-methylglutaryl-CoA synthase; JamI: enoyl hydratase/isomerase; JamJ: enoyl hydratase/isomerase/PKS protein; JamK: PKS protein; JamL: PKS/NRPS protein; JamM: PKS protein; JamO: NRPS protein; JamP: PKS/thioesterase; JamQ: cyclization.

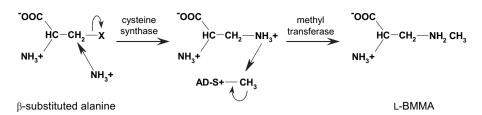


Fig. 7. Predicted two-step pathway for BMAA biosynthesis. From Brenner et al. (*Genome Biol.*, 2003, 4, R78, doi:10.1186/gb-2003-4-12-r78). X = phosphoserine, cysteine, *O*-acetylserine or cyanioalanine. Ad-S⁺ S-adenosyl-L-methionine.

L-BMAA (both, free and protein-bound forms; $8-287 \ \mu g \ g^{-1}$ cyanobacterial dry weight) and other known cyanobacterial toxins was found in bloom samples collected in British waterbodies between 1990 and 2004 (Metcalf et al., 2008).

Bioaccumulation of cyanobacterial L-BMAA could occur through different food chains in areas far from Guam and may explain the finding of L-BMAA in brain tissues of Alzheimer's patients from Canada (Cox et al., 2003; Murch et al., 2004a,b). Although there is mounting in vivo and in vitro evidence supporting a link between the presence of L-BMAA and ALS/PDC, the aetiology and pathogenesis of this neurodegenerative syndrome in the population of Western Pacific islands and in a small number of Caucasian and North American patients are still a matter of hot debate (Montine et al., 2005; Cox et al., 2006; Duncan and Marini, 2006; Garruto, 2006; Miller, 2006; Papapetropoulos, 2007; Steele and McGeer, 2008). Karymyan and Speth (2008) published a comprehensive review to which the reader is invited to refer for the description and critical evaluation of the animal models used for in vivo studies of L-BMAA neurotoxicity. Almost all these studies proved the neurotoxicity of this amino acid, which is generally associated with motor system disorder.

Based on analysis of expressed sequence tags, obtained from RNAs of young *Cycas rumphii* leaves, a two-step pathway has been proposed for the biosynthesis of L-BMAA in cycads (Brenner et al., 2003). In this pathway, L-BMAA synthesis begins with the transfer of NH₃ to a β -substituted alanine to form a metabolic intermediate. A cysteine synthase-like enzyme catalyzes this reaction. The second step is catalyzed by a methyl transferase that transfers a methyl group from S-adenosylmethionine to the new amine group of the intermediate compound (Fig. 7). Putative orthologous genes of both enzymes are present in cyanobacterial genomes, but gene inactivation is required to confirm that they are involved in the biosynthesis of L-BMAA in cyanobacteria (N. Tandeau de Marsac, unpublished data).

Considering the worldwide distribution of free-living and symbiotic cyanobacteria in most ecosystems, a search for the presence of L-BMAA in water supplies and human diets should be recommended. The ensemble of data collected, emphasizes the need for more research on L-BMAA and its relationships with neurodegenerative illnesses.

5. Conclusion

Cyanobacteria are ubiquitous primary producers that constitute up to 70% of the total phytoplankton biomass,

they produce more than 30% of total free O_2 and account for more than 30% of total primary production (CO₂ fixation). Cyanobacterial excessive proliferation is of serious concern for public health as they produce hepatotoxins and neurotoxins. However, the toxigenicity of cyanobacteria is not uniform and even at the species level there is a wide variation of virulence. Cyanobacterial neurotoxins described to date target cholinergic synapses and voltagegated sodium channels. The recent discovery of the lipopeptide neurotoxins antillatoxin, kalkitoxin and jamaicamide recalls the role of cyanobacteria as an important source for the discovery of emergent toxins. The complex structure of cyanobacterial neurotoxins represents a challenge for organic chemistry often leading to the development of novel synthetic pathways for their use as blueprints for the design of front-line drugs.

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Conflict of interest

The authors have no conflict of interest.

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