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Temporal variability of microcystin (mcyA) genotypes in a toxic cyanobacterial bloom.

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The cyanobacterium *Planktothrix agardhii* forms persistent blooms in water reservoirs and is often associated with the production of microcystins (hepatotoxins). However, high cell densities of P.agardhii are not always accompanied with high levels of microcystins and contrariwise. The genotype abundance and composition can be an important driver of toxicity in these natural blooms. In this work a perennial bloom was monitored during two years (2012-2014) in order to characterize the genotype structure and succession. A real-time PCR protocol was developed to quantify P.agardhii using the rpoC1 gene and to quantify potential microcystins producers using the microcystin syntethase gene (mcyA). The mcyA was quantified by targeting two gene regions: a generic region for all microcystin producers and a specific region for Planktothrix. Phytoplankton diversity and abundance was quantified by direct cell inspection and counting. The total microcystin concentration in water was measured using ADDA-ELISA kit. The results showed that two different mcyA genotypes are present in this water reservoir. The temporal variability accessed by cyanobacteria cell density, realtime PCR and microcystin concentration will be discussed. Furthermore, the real-time PCR protocol developed in this study enabled to determine and quantify genotype bloom composition, thus representing a promising tool in cyanobacteria bloom monitoring.