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## Noi siamo acqua

*Conoscere gli ecosistemi acquatici per riconnettersi alla natura* 





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## O.6.10 - Assessing the toxigenic potential of cyanobacteria in the Alpine region by combining high-throughput sequencing and metabolomic profiling

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In a recent survey carried out in the Alpine region as part of the Interreg Alpine Space Eco-AlpsWater project, we investigated the distribution of cyanobacteria and cyanotoxins in a large number of lakes. Plankton samples were collected monthly at 8 key lakes and 1 to 4 times at 30 additional sites in four countries. Taxonomic identifications and abundances of cyanobacteria were estimated by determining 16S rRNA amplicon sequence variants (ASVs) using highthroughput sequencing and light microscopy, while cyanotoxins were determined using liquid chromatography-mass spectrometry (LC-MS). Overall, in terms of relative abundance, the cyanobacteria showed a widespread presence of Chroococcales (mainly *Cyanobium*  sp.) and, especially in the largest water bodies, *Planktothrix rubescens*. In contrast, consistent pelagic populations of *Tychonema bourrellyi* were observed in the large lakes south of the Alps. The ASVs abundances of *P. rubescens* and *T. bourrellyi* showed a high correlation with microcystin and anatoxin-a concentrations, demonstrating a high consistency of the results obtained by HTS and metabolomic profiling. Overall, the results of the study showed a high capability of HTS to estimate the relative abundances of toxigenic cyanobacteria and their toxigenic potential.