## THE 18<sup>TH</sup> INTERNATIONAL CONFERENCE



FROM ECOSYSTEMS TO SOCIO-ECOSYSTEMS





## "Omic" technologies: genomics, transcriptomics, proteomics & metabolomics of HABs O-172

## Metabolic profiles of cyanobacteria isolated from Italian perialpine lakes

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**Abstract:** A comparative profiling of the secondary metabolism of cyanobacteria isolated from Italian perialpine lakes has been attempted. Cultures of *Aphanizomenon flos-aquae*, *Dolichospermum lemmermannii*, *Microcystis aeruginosa*, *Planktothrix rubescens*, and *Tychonema bourrellyi* were obtained from lakes Garda, Idro, and Caldonazzo. LC-MS/MS analysis were used for characterizing the strains' extracts. A target analysis aimed at assessing the the toxic profile (microcystins, nodularins, anatoxins, cylindrospermopsins, PSP) was performed; it revealed the production of toxic peptides (microcystins) in *M. aeruginosa* and *P. rubescens*, of toxic alkaloids (anatoxin-a and possibly some paralytic shellfish toxins) in *T. Bourrellyi*, and finally no production of toxins in *Aph. flos-aquae* and *D. Lemmermannii*. An untargeted analysis aimed at comparing the capability of the cyanobacterial species to produce peptidic metabolites, allowed the detection of over 300 different peptides, in a mass range between 400 and 2000 Da. The majority of compounds with masses between 500 and 1200 Da (corresponding to many non-ribosomal peptides) was produced by the two species *M. aeruginosa* and *P. rubescens*, which show a much higher capability of producing these compounds compared to the others. Microcystins, aeruginosins and anabaenopeptins were the most represented classes of compounds.

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