

SUPPLEMENTARY FIGURE 1

Phylogenetic relationships of cyanobacteria derived from neighbor-joining analysis of amino acid sequences of the core scytonemin genes *scyABC*. Numbers next to nodes represent non-parametric bootstrap support values based on 1000 replicates. Scytonemin-operon homologue sequences are demarcated by the orange bar, most closely related genes from other bacteria by the blue bar.

