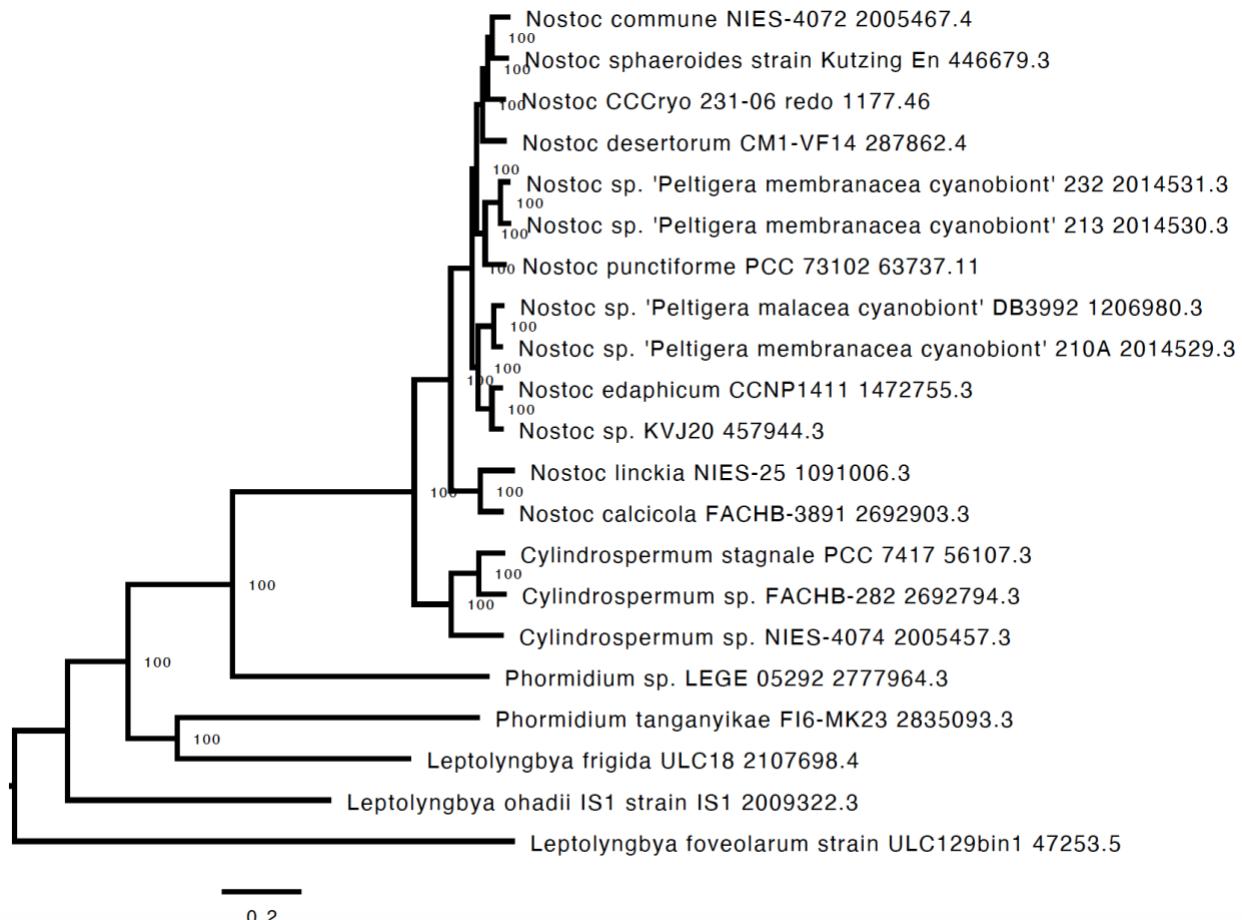


## **Supplemental information**

**Whole genome sequencing of cyanobacterium**

***Nostoc* sp. CCCryo 231-06 using microfluidic  
single cell technology**

**Yuguang Liu, Patricio Jeraldo, William Herbert, Samantha McDonough, Bruce Eckloff, Dirk Schulze-Makuch, Jean-Pierre de Vera, Charles Cockell, Thomas Leya, Mickael Baqué, Jin Jen, and Marina Walther-Antonio**



**Figure S1.** All 1000 genes in Nostoc – Related to Phylogenic tree building in Methods Details.