



Screening for Bio-H₂ production by cyanobacteria isolated from the Baltic Sea and Finnish lakes



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Cyanobacteria are capable of harnessing solar energy and converting it into chemical energy of carbohydrates in oxygenic photosynthesis. Under specific conditions, they can use solar energy to produce also molecular hydrogen. Cyanobacteria have three different enzymes that are directly involved in H₂ production.

Bidirectional hydrogenase (*hox*) has the ability to function both in uptake and in production of H₂. Bidirectional hydrogenases are pentameric, made of the hydrogenase moiety (HoxYH) and the diaphorase moiety (HoxEFU). The HoxEFU subunits are homologous to subunits of complex I of mitochondrial and bacterial respiratory chains. Uptake hydrogenases consisting of HupS and HupL subunits are present in all nitrogen fixing cyanobacteria strains. The uptake hydrogenase recycles the molecular hydrogen produced by nitrogenase in nitrogen fixation.

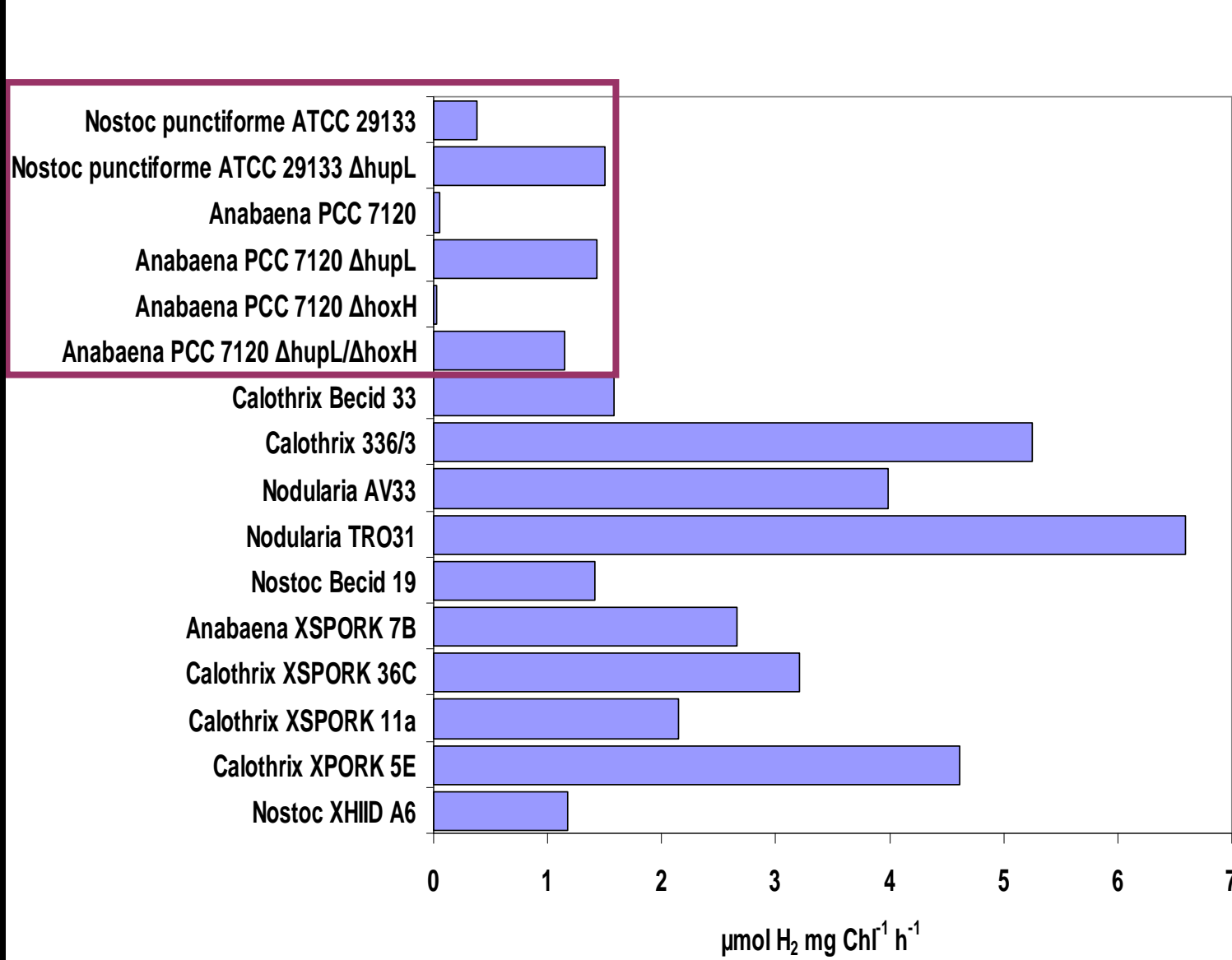
Nitrogenase produces hydrogen as a byproduct in nitrogen fixation. Mo-nitrogenase is made up of two parts, dinitrogenase consisting of two subunits each of NifD and NifK, and dinitrogenase

reductase, consisting of two subunits of NifH. *Anabaena variabilis* possesses two Mo-containing nitrogenases and also alternative vanadium(V)-containing nitrogenases (*vnf*).

Biodiversity among cyanobacteria for natural H₂ production has not been efficiently screened despite extensive research on H₂ production by microorganisms in general. Here we report the screening for H₂ production of 400 cyanobacterial strains isolated from the Baltic Sea and Finnish lakes. Approximately 50% of these strains produced detectable amounts of H₂. Each of the ten best H₂ producers belongs to N₂-fixing filamentous strains, eight of them are benthic and only two are planktonic strains.

The effect of different environmental factors, such as light intensity, cell density, pH and temperature on H₂ production rate was studied in the two good H₂ producers, *Calothrix* strains 336/3 and XPORK 5E. Notably, changing of environmental factors had differential effects on H₂ production depending on the strain.

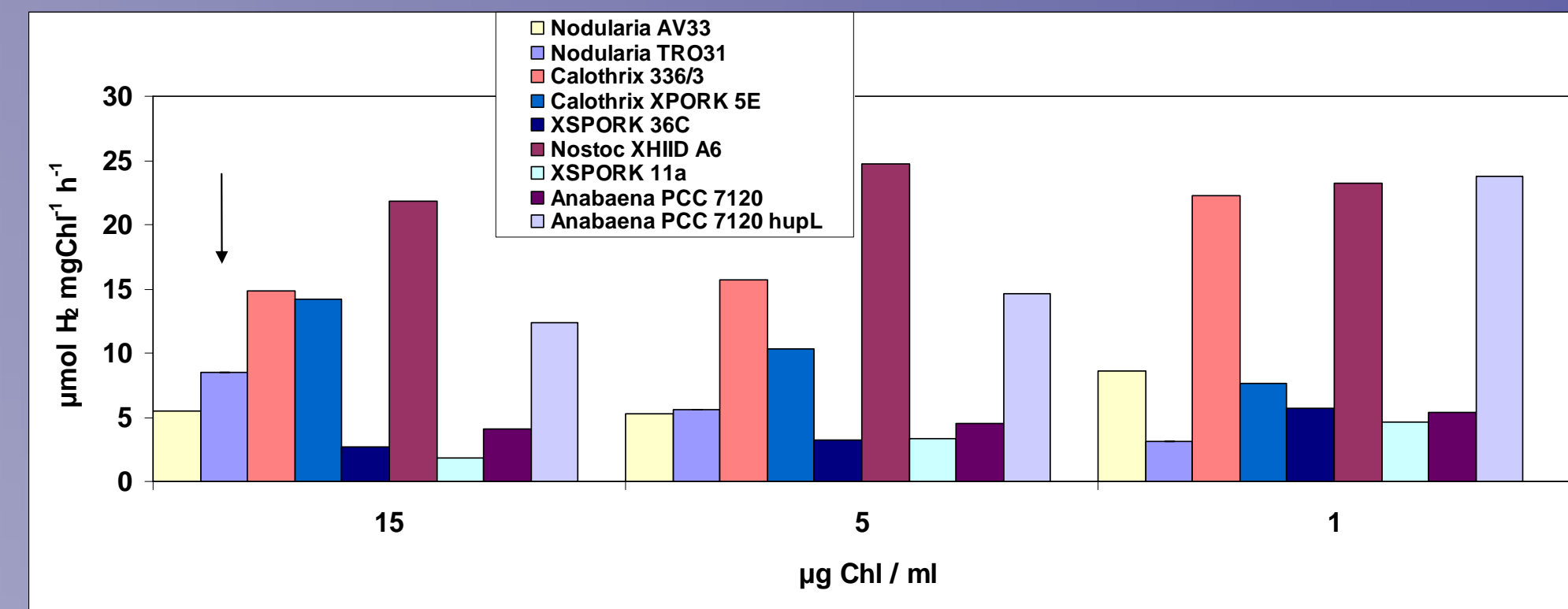
The rate of H₂ production of the ten best H₂ producers discovered during the screening process



Most of the H₂ producers were N₂-fixing filamentous strains, of 100 best producing strains 51 were planktonic, 46 benthic, and 3 soil/lichen associated. Only 1 unicellular and 1 non-N₂ fixing filamentous strain produced H₂. The highest H₂ production rates generally occurred under microaerobic/light conditions.

Figure demonstrates the rate of H₂ production of the 10 best H₂ producers discovered during the screening process. Uptake hydrogenase mutants of *Nostoc punctiforme* ATCC 29133⁽¹⁾ and *Anabaena PCC 7120*⁽²⁾ were used as positive controls in experiments.

The effect of decreasing cell density on the H₂ production rate in more optimized conditions



The H₂ production rate of most of the strains so far examined increases with lowering the cell density of the cultures. Nevertheless, some planktonic strains, like *Nodularia TRO31* demonstrated different behavior and lowering of the cell density of the cultures did not change or even lowered the H₂ production rate due to sensitivity to increased light intensity.

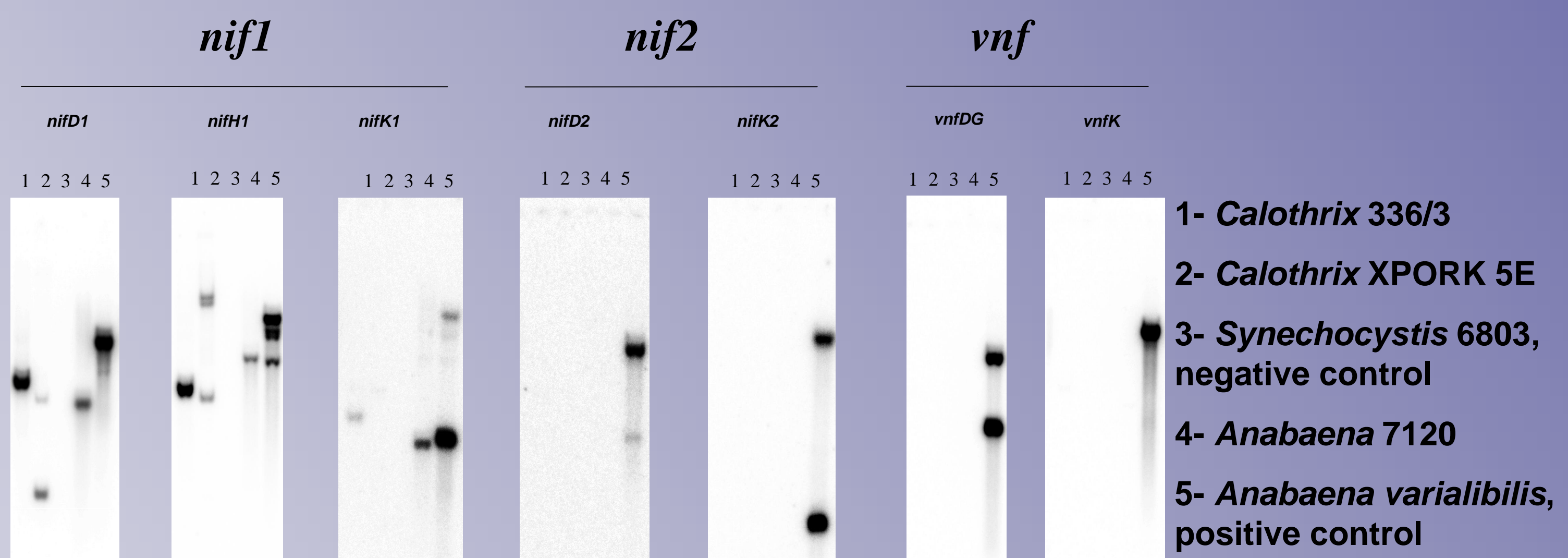
The origin of top 10 'superior' H₂ producers

Strain	Type	Origin	Habitat	Medium
<i>Calothrix</i> 336/3	filamentous	L. Enäjärvi, Laukilaanlahti,	benthos	Z8X
<i>Calothrix</i> BECID 33	filamentous	Gulf of Finland, the Baltic Sea	benthos	Z8X
<i>Nodularia</i> AV33	filamentous	Brackish water, the Baltic Sea	plankton	Z8XS
<i>Nodularia</i> TRO31	filamentous	Brackish water, the Baltic Sea	plankton	Z8XS
<i>Nostoc</i> BECID 19	filamentous	Vuosaari, the Baltic Sea coast	benthos	Z8X
<i>Nostoc</i> XHIID A6	filamentous	L. Hiidenvesi, Kirkkojärvi	benthos	Z8X
<i>Calothrix</i> XPORK 5E	filamentous	Porkkala cape, the Baltic Sea coast	benthos	Z8X
<i>Calothrix</i> XSPORK 11A	filamentous	Porkkala cape, the Baltic Sea coast	benthos	Z8XS
<i>Anabaena</i> XSPORK 36C	filamentous	Porkkala cape, the Baltic Sea coast	benthos	Z8XS
<i>Anabaena</i> XSPORK 7B	filamentous	Porkkala cape, the Baltic Sea coast	benthos	Z8XS

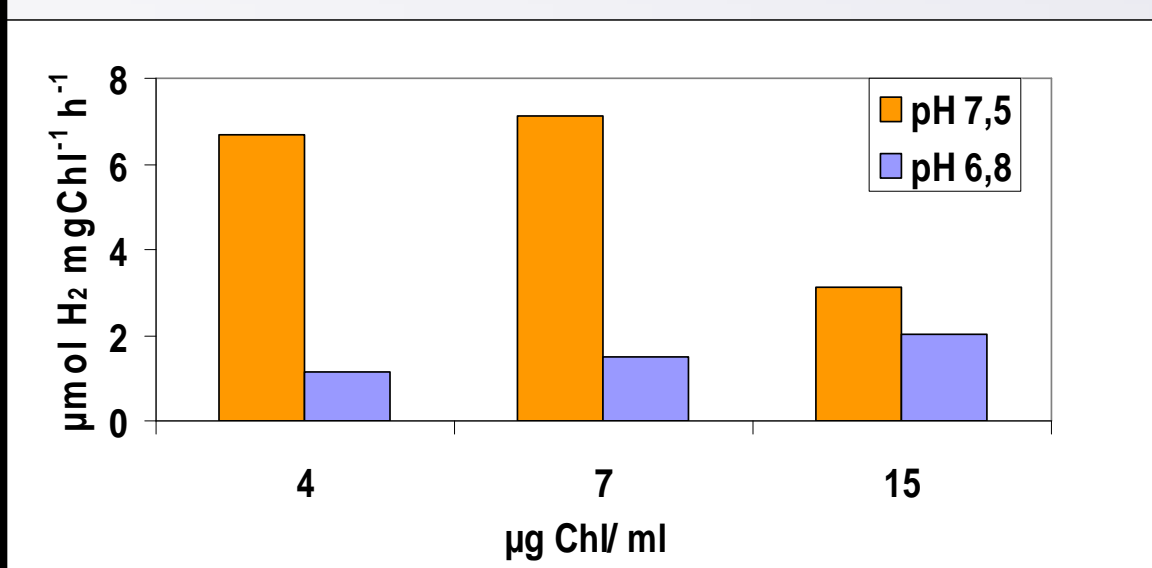


Detection of *nif1*, *nif2* and *vnf* genes by Southern hybridization

Southern hybridization of genomic DNA with *nif1*, *nif2* and *vnf* probes showed the presence of *nif1* Mo-nitrogenase in both *Calothrix* XPORK 5E and *Calothrix* 336/3, and absence of *nif2* and Vanadium nitrogenase in both strains.

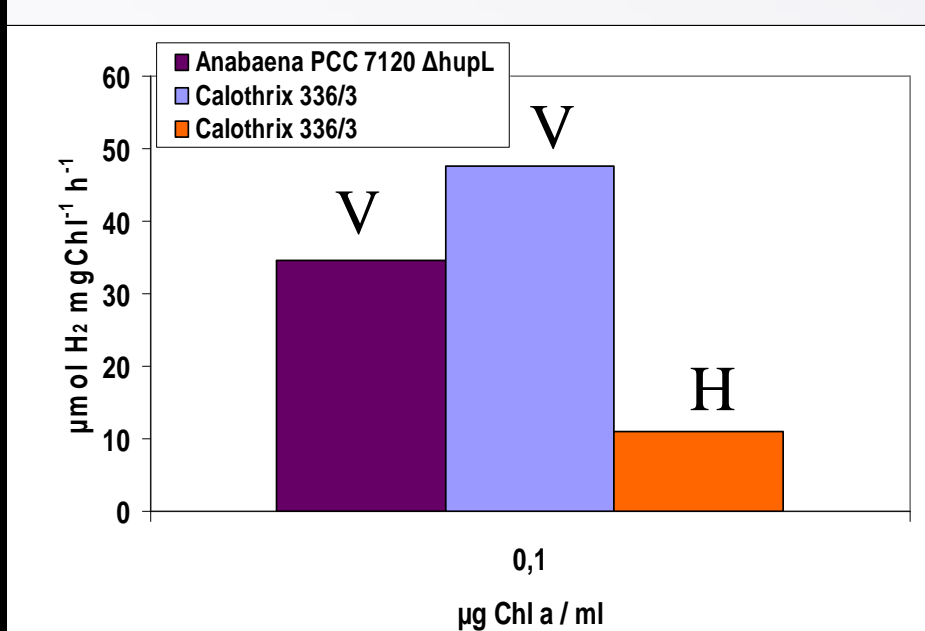


The effect of pH on the H₂ production rate of *Calothrix* XPORK 5E



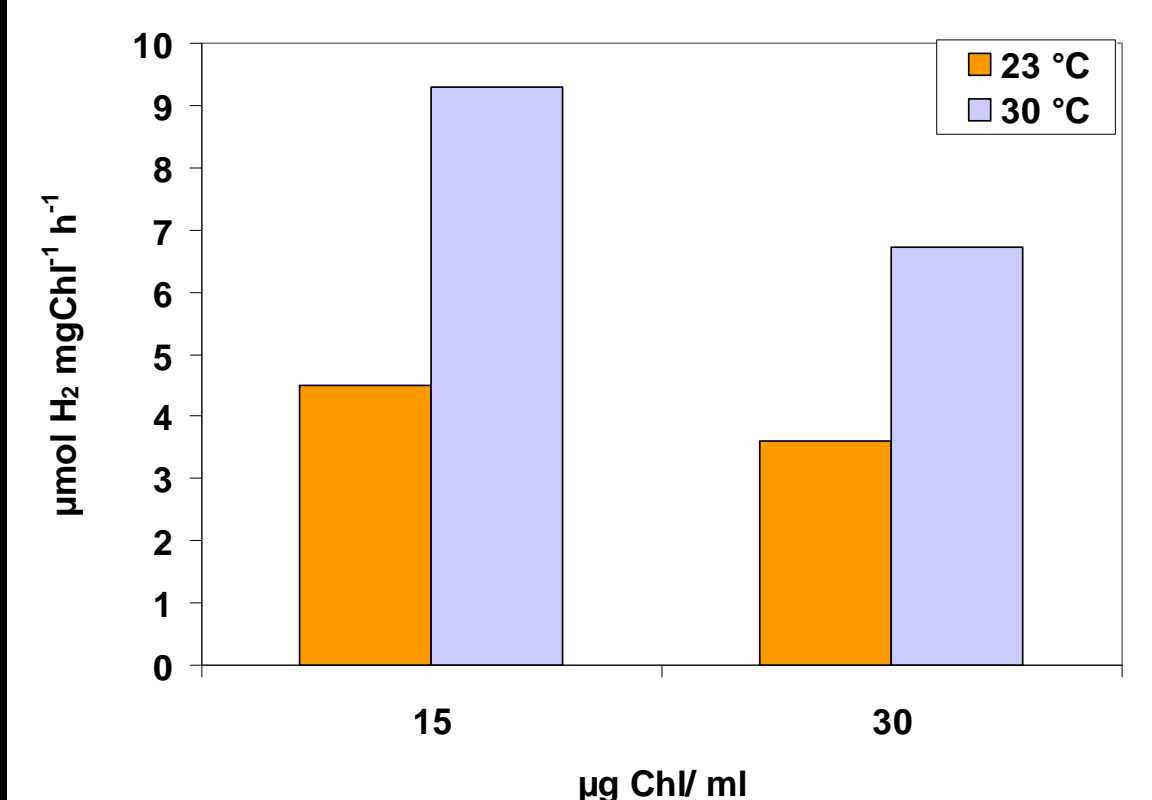
Calothrix XPORK 5E strain did not grow at pH 8.2 and grew relatively slowly at pH 6.8. At pH 6.8 *Calothrix* XPORK 5E produced much less H₂ compared to that at optimal pH 7.5.

The H₂ production rate of *Calothrix* 336/3 after optimizing the conditions



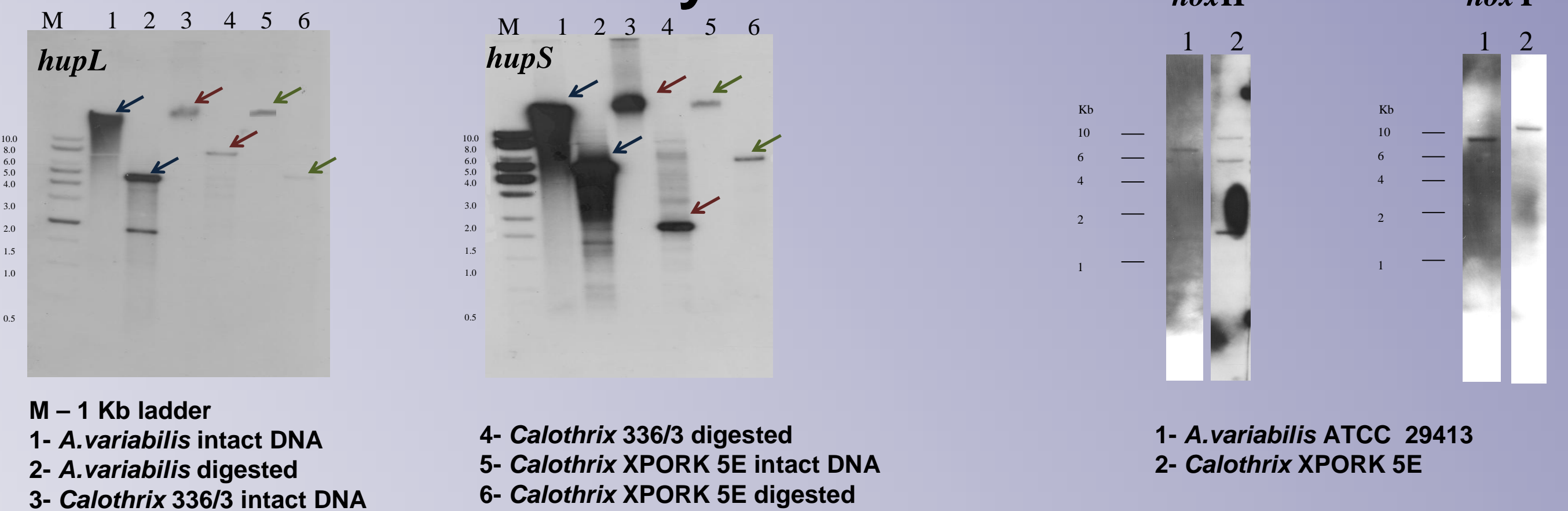
Calothrix 336/3 produced H₂ in much higher rate in vials incubated in horizontal position with larger surface area for light perception, more equal light distribution and more intensive mixing of the gas phase than in vials in vertical position.

The temperature effect on the H₂ production rate in *Calothrix* 336/3



By increasing the temperature of the H₂ determination assay from 23°C to 30°C the H₂ production rate of this strain was stimulated almost two fold.

Detection of *hup* and *hox* genes by Southern hybridization



Southern hybridization of genomic DNA with *hup* and *hox* probes showed the presence of uptake hydrogenase in both *Calothrix* 336/3 and *Calothrix* XPORK 5E, and the presence of bidirectional hydrogenase only in *Calothrix* XPORK 5E.

Future perspectives

Most of the superior strains with high H₂ production capacity, discovered by screening of the University of Helsinki Cyanobacteria Culture Collection (UHCC), are benthic microorganisms with high capacity to adhere to the solid substrates. In collaboration with Dr. Tsygankov's group (Institute of Basic Biological Problems, Pushchino, Russia) we will apply immobilization and continues culture techniques to these strains in order to further improve their H₂ production activity.

We are taking a systems biology approach to characterise these strains in detail for maximising the H₂ producing capacity. The sequencing of the genomes of the two novel 'superior' strains, will be performed by submission them to the next-generation sequencing system SOLiD™ in collaboration with researchers in the Center of Biotechnology (BTK), Biocity, Turku. In long run, this will give us an opportunity to further improve the H₂ production capacity of these strains by genetic engineering techniques.

Ref: 1) Lindberg P., Schutz K., Happe T. & Lindblad P. Int. J. Hydrogen Energy 2002; 27, 1291-1296
2) Masukawa H., Mochimaru M. & Sakurai H. App. Microbiol. Biotech. 2002; 58, 618-624