## Isolation of the first thermophilic and actively nitrogen-fixing bacteria in the deep branching phylum *Aquificae*

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Thermophilic nitrogen-fixing bacteria have been suggested to occur in hydrothermal vents and terrestrial hot springs from molecular and functional-based analyses of prokaryotic communities. Nif genes, biomarkers for nitrogen-fixing bacteria, already have been found in some chemosynthetic thermophilic isolates, too. Although nitrogen-fixing ability was observed in methanogenic archaea at 92°C, active nitrogen fixation in thermophilic bacteria more than 70°C has not been demonstrated yet. In this study, we isolated two novel Aquificae, a deeply branching bacterial phylum, into axenic culture under nitrogen-fixing conditions from chemosynthetic microbial communities at 70-77°C in sulfidic alkaline hot springs (Nakabusa, Nagano, Japan). Phylogenetic analysis based on 16S rRNA gene classified both strains within the genus Hydrogenobacter, strain 1-6 showed 98.7% nt identity to Hydrogenobacter subterraneus HGP1 and strain 2-18 had 97.6% identity to H. hydrogenophilus DSM 2913. Both isolated strains contained nifH gene sequences, encoding a key enzyme component of nitrogen fixation, with 96.5% and 97.4% amino acid identity to Hydrogenobacter thermophilus TK-6. Nitrogenase activities were confirmed in both strains incubated at 70°C using the acetylene reduction test. Both strains showed nitrogen-gas-dependent growth under lowered aerobic conditions with approximately up to 10% oxygen using CO<sub>2</sub> as a sole carbon source and N<sub>2</sub> as sole nitrogen source with H<sub>2</sub> or thiosulfate as electron donors at 70°C.

This is the first demonstration of active nitrogen-fixation in thermophilic bacteria (more than 70°C) and in the phylum *Aquificae*. The potential impact of nitrogen fixation in thermophilic chemosynthetic bacteria will be discussed with respect to life on earth before the appearance of photosynthesis.

Keywords: Aquificales, nitrogen fixation, geothermal spring, thermophile, chemosynthetic bacteria