

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

\*西原 亜理沙<sup>1</sup>、松浦 克美<sup>1</sup>、Tank Marcus<sup>1</sup>、McGlynn Shawn<sup>2,3,4</sup>、Thiel Vera<sup>1</sup>、春田 伸<sup>1</sup>

\*Arisa Nishihara<sup>1</sup>, Katsumi Matsuura<sup>1</sup>, Marcus Tank<sup>1</sup>, Shawn E McGlynn<sup>2,3,4</sup>, Vera Thiel<sup>1</sup>, Shin Haruta<sup>1</sup>

1. 首都大学東京、2. 東京工業大学 地球生命研究所、3. 理化学研究所 環境資源科学研究センター、4. Blue Marble Space Institute of Science

1. Tokyo Metropolitan University, 2. Earth-Life Science Institute, Tokyo Institute of Technology, 3. Biofunctional Catalyst Research Team, RIKEN Center for Sustainable Resource Science, 4. Blue Marble Space Institute of Science

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.

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