Geoglobus acetivorans sp. nov., an iron(III)-reducing archaeon from a deep-sea hydrothermal vent

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Abstract:

A hyperthermophilic, anaerobic, dissimilatory Fe(III)-reducing, facultatively chemolithoautotrophic archaeon (strain SBH6T) was isolated from a hydrothermal sample collected from the deepest of the known World Ocean hydrothermal fields, Ashadze field (1 ° 58' 21'' N 4 ° 51' 47'' W) on the Mid-Atlantic Ridge, at a depth of 4100 m. The strain was enriched using acetate as the electron donor and Fe(III) oxide as the electron acceptor. Cells of strain SBH6T were irregular cocci, 0.3–0.5 µm in diameter. The temperature range for growth was 50–85 °C, with an optimum at 81 °C. The pH range for growth was 5.0–7.5, with an optimum at pH 6.8. Growth of SBH6T was observed at NaCl concentrations ranging from 1 to 6 % (w/v) with an optimum at 2.5 % (w/v). The isolate utilized acetate, formate, pyruvate, fumarate, malate, propionate, butyrate, succinate, glycerol, stearate, palmitate, peptone and yeast extract as electron donors for Fe(III) reduction. It was also capable of growth with H2 as the sole electron donor, CO2 as a carbon source and Fe(III) as an electron acceptor without the need for organic substances. Fe(III) [in the form of poorly crystalline Fe(III) oxide or Fe(III) citrate] was the only electron acceptor that supported growth. 16S rRNA gene sequence analysis revealed that the closest relative of the isolated organism was Geoglobus ahangari 234T (97.0 %). On the basis of its physiological properties and phylogenetic analyses, the isolate is considered to represent a novel species, for which the name Geoglobus acetivorans sp. nov. is proposed. The type strain is SBH6T (=DSM 21716T =VKM B-2522T).

Abbreviations: AQDS, 9,10-anthraquinone-2,6-disulfonate; DGGE, denaturing gradient gel electrophoresis
Iron minerals are abundant in the deep-sea hydrothermal vents. The surfaces of active chimneys are frequently covered with deposits of iron oxides at different oxidative states and amount of iron in hydrothermal fluid can reach molar concentrations. Thus, deep-sea hydrothermal vents can provide ecological niche for Fe(III)-reducing micro-organisms (Slobodkin, 2001). However, only few thermophilic Fe(III)-reducers were isolated from this environment. Currently, thermophilic and hyperthermophilic iron-reducing micro-organisms recovered from deep-sea habitats include two species of Bacteria, Geothermobacter ehrlichii (Kashefi et al., 2003) and Deferribacter abyssii (Miroshnichenko et al., 2003) and three representatives of Archaea, Thermococcus sp. SN531 (Slobodkin et al., 2001), Geoglobus ahangari (Kashefi et al., 2002) and Aciduliprofundum boonei (Reysenbach et al., 2006).

In this article, we report the isolation and characterization of the novel hyperthermophilic Fe(III)-reducing archaeon from the deepest of the known World Ocean hydrothermal field.

Strain SBH6T was isolated from a sample of the fragment of the hydrothermal chimney-like structure. The sample was collected in March 2007 during the SERPENTINE cruise at the Ashadze hydrothermal field (12 58’ 21” N, 44 51’ 47” W) on the Mid-Atlantic Ridge at a depth of 4100 m. For the samples collection sterilized microbiological boxes filled with sterile freshwater were prepared onboard. Active chimney samples were collected by the ROV Victor. On site, after opening the box lid, the freshwater was replaced by in situ seawater, the chimney fragment introduced and the lid closed. All following operations were done onboard in sterile conditions. Boxes with samples were stored at 4°C. An enrichment culture was initiated by inoculation of 10% (w/v) of the sample into anaerobically prepared, bicarbonate-buffered, sterile (135°C, 1 h) liquid medium with acetate (18 mM) as an electron donor and poorly crystalline Fe(III) oxide (90 mmol/l⁻¹) as an electron acceptor. Medium composition and preparation techniques were described earlier (Slobodkin et al., 1999a); the medium was additionally supplemented with NaCl (18 g l⁻¹) and MgCl₂ (4 g l⁻¹) to increase the salinity. After three subsequent transfers and following serial 10-fold dilutions in the same medium at 65°C two morphological types of cells were observed in the highest positive dilution (10⁻⁵): rods and cocci. To obtain coccoid-shaped micro-organism further cultivation was carried out at 82°C. After two transfers at 82°C the rods were not observed neither at 82°C nor at 65°C and only coccoid-shaped cells were present in the enrichment. Acetate (4.5mM) consumption in enrichments measured by gas chromatography reached 80%. The attempts to obtain separate colonies in agar-blocks or by roll-tube method (Hungate, 1969) were unsuccessful either at 82°C or at 65°C with 1% of GELRITE gellan gum or with 2% of agar as solidifying agent in the medium correspondingly. Then the enrichment was subsequently transferred five times in lithoautotrophic conditions with molecular hydrogen as an electron donor, poorly crystalline Fe(III) as the electron acceptor and CO₂ as the carbon source at 82°C and after that serially diluted at the same conditions. The isolate from the highest dilution that exhibited Fe(III) reduction (10⁻⁶), was considered as a pure culture and was designated strain SBH6T. Denaturing gradient gel-electrophoresis (DGGE) of the strain SBH6T grown on the media containing acetate (18 mM) or peptone (10 g l⁻¹) as electron donor and poorly crystalline Fe(III) oxide (90 mmol/l⁻¹) as electron acceptor in the presence of yeast extract (0.02 g l⁻¹) revealed the single band in each case and thus confirmed the purity of the culture. Complete 16S rRNA gene sequences of the strain SBH6T grown with acetate or peptone were identical. Physiological studies on substrate and electron acceptor utilization, temperature, pH and salinity ranges for growth, light and electron microscopy, analytical techniques (Fe(II) and acetate concentrations), DNA extraction were performed as described previously (Slobodkin et al., 1999a). Growth of the strain with poorly crystalline Fe(III) was determined by direct cells count using light microscopy after dissolving the iron precipitate in solution of ammonium oxalate (28 g l⁻¹) / oxalic acid (15 g l⁻¹) (Lovley and Phillips, 1988). pH measurements and pH-meter calibration were carried out at 60°C. 16S rRNA gene amplification, sequencing and sequence analysis were done as described previously (Zavarzina et al., 2002). 16S rRNA gene fragments for DGGE were obtained using PCR with primers Uni515F (Lane, 1991) withGC-clamp (Muyzer et al., 1993) at the 5’ end and 915R (Casamayor et al., 2002). DGGE was performed as described (Muyzer et al., 1997) with
denaturating gradient ranging from 35% to 65% (100% denaturant contains 7 M urea and 40% formamide).

Cells of strain SBH6<sup>T</sup> were regular to irregular cocci, about 0.3-0.5 μm in diameter, usually arranged as single cells, flagella were not observed. The temperature range for growth of strain SBH6<sup>T</sup> was 50-85°C, with an optimum at 81°C. No growth was detected at 90°C or at 46°C after incubation for 3 weeks. The pH range for growth was pH 5.0-7.5, with an optimum at pH 6.8. No growth was noticed at pH 4.5 or 8.0. Growth of strain SBH6<sup>T</sup> was observed at NaCl concentrations ranging from 1.0 to 6.0% (w/v) with an optimum at 2.5% (w/v), but no growth was evident at 0 and 7.0% NaCl (w/v). Poorly crystalline Fe(III) oxide was reduced to black magnetic precipitate with high Fe(II) content. No changes in color and amount of precipitate were observed in uninoculated controls containing poorly crystalline Fe(III) oxide during the incubation period at 82°C. Strain SBH6<sup>T</sup> grew and reduced Fe(III) with acetate (4.5 or 18 mM), formate, pyruvate, fumarate, malate, propionate, butyrate, succinate, glycerol, (20 mM each), stearate, palmitate (0.5-1.0 mM each), peptone, yeast extract, (10 g l<sup>−1</sup> each). During growth with acetate (4.5 mM) and Fe(III) in the absence of yeast extract, 80% of acetate were consumed and a ratio of Fe(II) produced to acetate consumed was 7.5. Strain SBH6<sup>T</sup> also grew with molecular hydrogen as the sole electron donor for Fe(III) reduction and CO<sub>2</sub> as the carbon source. No organic carbon source was required for growth on hydrogen. There was no Fe(III) reduction or cell growth in the absence of added hydrogen. Strain SBH6<sup>T</sup> was not able to utilize lactate (25mM), L-alanine, glycine (20 mM each), L-proline, arginine, serine, glutamine, asparagine, L-cysteine, aspartic acid, glutamic acid (10 mM each), methanol, ethanol, benzoate (20 mM each) with poorly crystalline Fe(III) oxide as an electron acceptor. Besides poorly crystalline Fe(III) oxide, strain SBH6<sup>T</sup> also could grow with Fe(III) citrate (10 mM) as the electron acceptor but cells yield and Fe(III) reduction was lower. Fe(III) citrate was not completely reduced, no more than 5-6 mM of Fe(II) was formed. Several attempts to grow strain SBH6<sup>T</sup> on a variety of commonly considered electron acceptors (including sulfate (14 mM), thiosulfate (20 mM), elemental sulfur (10 g l<sup>−1</sup>), nitrate (10 mM), fumarate (20 mM), Mn(IV)oxide (25 mM), 9,10-anthraquinone 2,6-disulfonate (AQDS (20 mM)) and oxygen (2 or 20%, v/v)) other than poorly crystalline Fe (III) oxide (90 mmol/l<sup>−1</sup>) and Fe(III) citrate (10 mM) using acetate (18mM), H<sub>2</sub> (as H<sub>2</sub>:CO<sub>2</sub>, 80:20%, v/v, 101 kPa), lactate (25 mM), butyrate (20mM) or glycerol (20 mM) were unsuccessful.

A comparison of 1417 nucleotides of 16S rRNA gene sequence of strain SBH6<sup>T</sup> with those available in GenBank database showed that strain SBH6<sup>T</sup> had the highest identity with *Geoglobus ahangari* 234<sup>T</sup> (97.0%) (Fig.1). Only 16S rRNA sequences of the type strains of validly published species were included in the analyses. The levels of 16S rRNA gene sequence similarity with other members of the order *Archaeoglobales* were 94.9-95.3%. The trees constructed by maximum likelihood and by maximum parsimony algorithms displayed the same topology (data not shown). Transversion analysis (Woese et al., 1991) did not affect the phylogenetic position of the new strain.

The new hyperthermophilic isolate described in this report, capable of reduction of Fe(III) was recovered from the environmental sample using acetate as the electron donor for initial enrichment. Acetate is one of the major metabolic products of organic matter decomposition under anaerobic conditions and it could be produced during fermentation by many hyperthermophiles (Slobodkin et al., 1999b). For a long time there was no data demonstrating anaerobic acetate degradation by hyperthermophiles. Then this ability was shown for two micro-organisms, *Ferroglobus placidus* and *Geoglobus ahangari* (Tor et al., 2001), however none of these strains was initially enriched and obtained into pure culture with acetate as the electron donor. Final purification of the isolate SBH6 was carried out in lithoautotrophic conditions with molecular hydrogen as the electron donor since colonies on the medium with acetate were not formed. However, already in enrichments cultivated in hyperthermophilic conditions only one morphological type of cells was observed. Since the same extent of acetate consumption and Fe(III) reduction was detected before and after...
purification in autotrophic conditions, we can assume that strain SBH6 was responsible for acetate utilization in initial enrichments. Therefore, SBH6 is the first hyperthermophilic micro-organism enriched on acetate as the electron donor. At the present, the order Archaeoglobales includes one family Archaeoglobaceae consisting of three genera: Archaeoglobus, Ferroglobus and Geoglobus (Cole et al., 2007). The genus Geoglobus is represented by the sole species, Geoglobus ahangari (Kashefi et al., 2002) isolated from a deep-sea hydrothermal sample from Guaymas Basin, Gulf of California. The isolate SBH6\textsuperscript{T} shares many phenotypic features with the described representative of this genus. First of all, it is the inability to use for growth the electron acceptors other than Fe(III), preferable insoluble Fe(III) oxide. Both micro-organisms grow poorly in media with soluble form of Fe(III) (Fe(III) citrate) as an electron acceptor. There is practically no difference in electron donors utilization, including anaerobic oxidation of long-chain fatty acids and the ability to grow chemolithoautotrophically. Significant differences between G. ahangari and the strain SBH6\textsuperscript{T} are noticed in growth temperatures. Strain SBH6\textsuperscript{T} grows at 50°C and it does not grow at 85°C and above whereas the low growth limit for G. ahangari is 65°C and it is able to grow up to 90°C. The optimal growth temperatures also differ for 7°C being 88 °C for G. ahangari and 81°C for the strain SBH6\textsuperscript{T}. In addition, analysis of 16S rRNA gene sequences revealed considerable phylogenetic distance between G. ahangari and isolate SBH6\textsuperscript{T}. Thus, phylogenetic and physiological properties clearly differentiate strain SBH6\textsuperscript{T} from the closest relative, G. ahangari. According to the opinion of Judicial Commission of the International Committee for Systematics of Prokaryotes (Judicial Commission of the International Committee for Systematics of Prokaryotes, 2008) G. ahangari is not validly published since it is deposited only in one collection of micro-organisms. However, we hope that it will be validated and we propose strain SBH6\textsuperscript{T} as the type strain of the new species, Geoglobus acetivorans, the second species of the genus Geoglobus.

**Description of Geoglobus acetivorans sp.nov.**

*Geoglobus acetivorans* (a.ce.ti.vo'rans. L. neut. n. acetum, vinegar or acetic acid; L. part. adj. vorans devouring; N.L. part. adj. acetivorans, vinegar consuming). Cells are regular to irregular cocci, 0.3-0.5 μm in diameter, occurring singly. The temperature range for growth is 50-85°C, with an optimum at 81°C. The pH range for growth is 5.0-7.5, with an optimum at pH 6.8. Growth occurs at NaCl concentrations ranging from 1.0 to 6.0% (w/v) with an optimum at 2.5% (w/v). Anaerobic. Capable of chemolithoautotrophic growth using molecular hydrogen as an electron donor, ferric iron as electron acceptor and CO\textsubscript{2} as the carbon source. Only poorly crystalline Fe(III) oxide and Fe(III) citrate are used as electron acceptors for growth. Sulfate, thiosulfate, elemental sulfur, nitrate, fumarate, Mn(IV)oxide, 9,10-anthraquinone 2,6-disulfonate and oxygen are not utilized as electron acceptors. With poorly crystalline Fe(III) oxide anaerobically oxidizes acetate, formate, pyruvate, fumarate, malate, propionate, butyrate, succinate, glycerol, stearate, palmitate peptone and yeast extract. Lactate, L-alanine, glycine, L-proline, arginine, serine, glutamine, asparagine, L-cysteine, aspartic acid, glutamic acid, methanol, ethanol, benzoate are not utilized. Isolated from deep-sea hydrothermal field (Ashadze) of the Mid-Atlantic Ridge. The type strain is SBH6\textsuperscript{T} has been deposited in the Deutsche Sammlung von Mikroorganismen und Zellkulturen under the accession number DSM 21716\textsuperscript{T} and in the All-Russian National Collection of Microorganisms (VKM) under the accession number VKM-2522\textsuperscript{T}. 

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References


Fig. 1. Phylogenetic tree based on 16S rRNA gene sequences indicating the position of isolate SBH6\(^T\) within the representative members of the order Archaeoglobales. The 16S rRNA gene sequence of *Thermoproteus tenax* was included as outgroup. GenBank accession numbers are given in parentheses. Bar, 10 substitutions per 100 nt. Only the bootstrap values higher than 70% are indicated.

- *Thermoproteus tenax* DSM 2078\(^T\) (M35966)
- *Thermococcus celer* DSM 2476\(^T\) (M21529)
- *Ferroglobus placidus* DSM 10642\(^T\) (AF220166)
- *Archaeoglobus lithotrophicus* TF2\(^T\) (AJ299218)
- *Archaeoglobus fulgidus* VC-16\(^T\) (X05567)
- *Geoglobus acetivorans* SBH6\(^T\)
- *Archaeoglobus profundus* DSM 5631\(^T\)
- *Archeoglobus infectus* Arc51\(^T\) (AB274307)
- *Archeoglobus veneficus* DSM 11195\(^T\) (AF418181)
- *Archaeoglobus profundus* DSM 5631\(^T\)
- *Ferroglobus placidus* DSM 10642\(^T\) (AF220166)
- *Thermococcus celer* DSM 2476\(^T\) (M21529)
- *Geoglobus ahangari* 234\(^T\) (AF220165)