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## ***Endothiovibrio diazotrophicus* gen. nov., sp. nov., a novel nitrogen-fixing, sulfur-oxidizing gammaproteobacterium isolated from a salt marsh**

Dennis A Bazylnski<sup>1</sup>, Viviana Morillo<sup>1</sup>, Christopher T Lefèvre<sup>2</sup>, Nathan Vilorio<sup>1</sup>, Bradley L Dubbels<sup>3</sup>, Timothy J Williams<sup>4</sup>

1 School of Life Sciences, University of Nevada at Las Vegas, Las Vegas, Nevada 89154-4004, USA

2 CEA Cadarache/CNRS/Université Aix-Marseille, UMR7265 Biosciences and Biotechnologies Institute, Laboratoire de Bioénergétique Cellulaire, Saint Paul lez Durance 13108, France

3 Novozymes North America Inc., 9000 Development Drive, Morrisville, North Carolina 27560, USA

4 School of Biotechnology and Biomolecular Sciences, The University of New South Wales, Sydney NSW 2052, Australia

\*Correspondence: Dennis A. Bazylnski, dennis.bazylnski@unlv.edu

### **ABSTRACT**

A novel non-phototrophic, marine, sulfur-oxidizing bacterium, strain S-1<sup>T</sup>, was isolated from a coastal salt marsh in Massachusetts, USA. Cells are Gram-stain-negative vibrios motile by means of a single polar unsheathed flagellum. S-1<sup>T</sup> is an obligate microaerophile with limited metabolic capacity. It grows chemolithoautotrophically utilizing sulfide and thiosulfate as electron donors, converting these compounds to sulfate, and the Calvin–Benson–Bassham cycle for carbon fixation. Cells of S-1<sup>T</sup> did not grow on any of a large number of organic carbon sources and there was no evidence for chemoorganoheterotrophic growth. Cells produced internal sulfur globules during growth on sulfide and thiosulfate. S-1<sup>T</sup> is strongly diazotrophic, as demonstrated by <sup>15</sup>N<sub>2</sub> fixation and acetylene reduction activity by cells when a fixed nitrogen source is absent from the growth medium. The marine nature of this organism is evident from its ability to grow in 10 to 100% artificial seawater but not at lower concentrations and NaCl alone cannot substitute for sea salts. The major cellular fatty acids are C<sub>16:1</sub>ω7c, C<sub>16:0</sub>, and C<sub>18:1</sub>ω7c. Phosphatidylethanolamine and phosphatidylglycerol are the major polar lipids. Q8 is the only respiratory quinone. S-1<sup>T</sup> genomic DNA has a G+C content of 67.6 mol%. Based on its 16S rRNA gene sequence, S-1<sup>T</sup> shows the closest phylogenetic relationship to non-phototrophic species within the family *Thioalkalispiraceae* of the class *Gammaproteobacteria*. The name *Endothiovibrio diazotrophicus* is proposed for this organism, with S-1<sup>T</sup> as the type strain (ATCC BAA-1439<sup>T</sup>=JCM 17961<sup>T</sup>).

**Keyword(s):** sulfur deposits, sulfur oxidization, Thioalkalispiraceae, S-1, nitrogen fixation, *Endothiovibrio diazotrophicus*

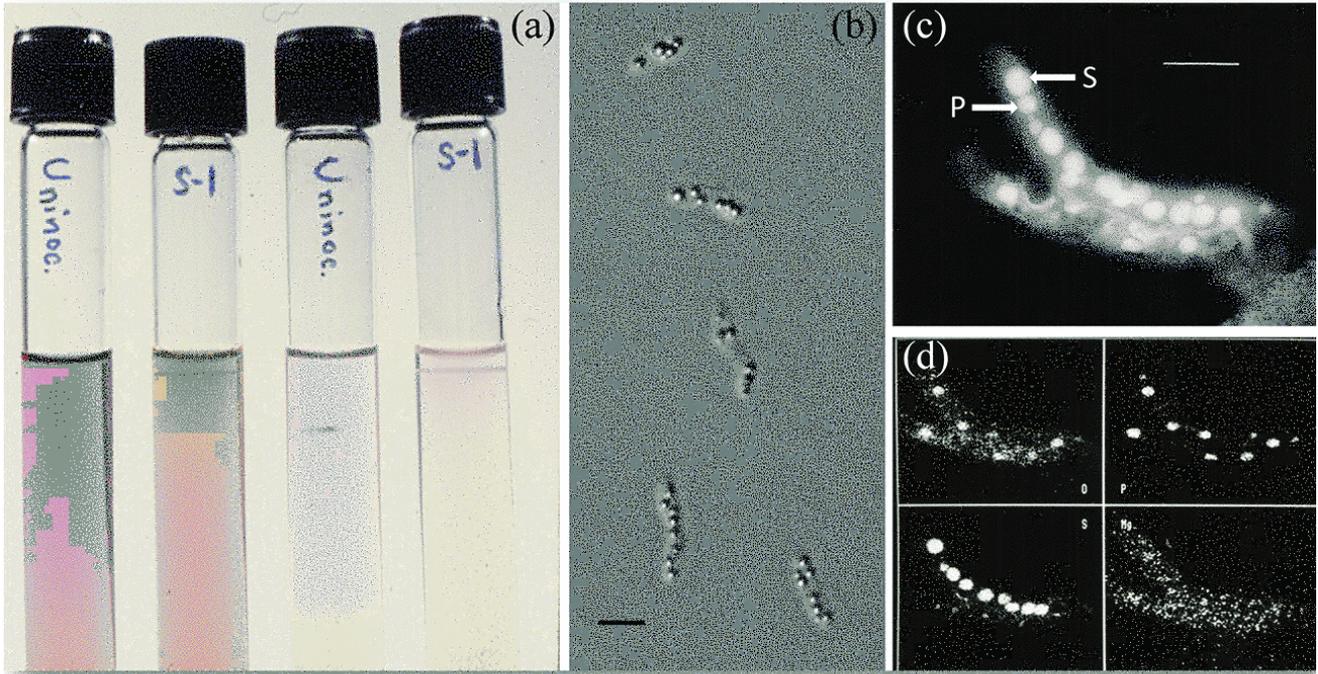
The *Gammaproteobacteria* class of the phylum *Proteobacteria* is one of the most genera-rich taxa among the bacteria [ 1 ] and includes a relatively large number of chemolithoautotrophic and photolithoautotrophic sulfur-oxidizing bacteria. Most of these belong to one of two orders: the *Chromatiales* (the ‘purple sulfur bacteria’) and the *Thiotrichales*. Although the ability to store and metabolize elemental sulfur is widespread among these bacteria, only a relatively small subset is known to store sulfur intracellularly [ 2 ]. In these cells, sulfur inclusions are stored within the confines of the cell wall, and have an extracytoplasmic/periplasmic location [ 2, 3 ]. In many chemotrophic and phototrophic sulfur-oxidizers, the deposition of elemental sulfur appears to be an obligatory intermediate step in the complete oxidation of sulfide and thiosulfate to sulfate, probably for kinetic reasons [ 2, 4 ]. The formation of these so-called sulfur ‘globules’ is prevalent among

members of the order *Chromatiales* of the class *Gammaproteobacteria*, in which the oxidation of reduced sulfur compounds fuels the autotrophic assimilation of CO<sub>2</sub> (thioautotrophy). The order *Chromatiales* can be classified into seven families: *Chromatiaceae*, *Ectothiorhodospiraceae*, *Halothiobacillaceae*, *Thioalkalispiraceae*, *Wenzhouxiangellaceae*, *Woeseiaceae* and *Granulosicoccaceae* [ 5–8 ]. Here we describe strain S-1<sup>T</sup>, a marine, non-phototrophic, sulfur-oxidizing member of the family *Thioalkalispiraceae*, a family of strict chemolithoautotrophs that are mesophilic/moderately thermophilic and moderately halophilic [ 6 ]. S-1<sup>T</sup> exhibits intracellular sulfur globule deposition associated with the oxidation of sulfide and thiosulfate.

Mud and water was collected from the brackish School Street Marsh at Woods Hole, (Massachusetts, USA) and used to inoculate sulfide-O<sub>2</sub> concentration gradient media, prepared following the recipe of Nelson and Jannasch [ 9 ] but modified by using a diluted artificial seawater (ASW) solution to replace natural seawater and by the addition of 20 mM ferric quinate and 200 µl 0.2% aqueous resazurin per litre. The ferric quinate was originally included in the growth medium because we were attempting to culture magnetotactic bacteria and later removed because the organism described here did not require this high concentration of iron for growth. The salinity of the water at the site was 10 p.p.t. (=1% w/v) as determined with a Palm Abbe PA203 hand-held refractometer (MISCO Refractometer). The ASW used in routine culturing of this organism was adjusted to this salinity and consisted of (g l<sup>-1</sup>): NaCl, 7.82; MgCl<sub>2</sub>·6H<sub>2</sub>O, 1.66; Na<sub>2</sub>SO<sub>4</sub>, 1.30; KCl, 0.22; and CaCl<sub>2</sub>·2H<sub>2</sub>O, 0.18. Cells initially grew at the oxic–anoxic interface (OAI), corresponding to the pink–colorless interface as a microaerobic band of cells. Although the culture was mixed, the dominant microorganism was a vibrio that appeared to produce intracellular sulfur globules.

For isolation of the strain, cells from the enrichment gradient culture were inoculated in a dilution series of solid agar [13 g l<sup>-1</sup> Agar Noble (Difco Laboratories)] in shake tubes of an O<sub>2</sub> concentration gradient medium containing: 5 ml modified Wolfe's mineral elixir [ 10 ]; 0.5 ml vitamin solution [ 10 ]; 10 mM Na<sub>2</sub>S<sub>2</sub>O<sub>3</sub>·5H<sub>2</sub>O as the electron donor; 0.2 g NH<sub>4</sub>Cl as the nitrogen source; 2 ml freshly prepared, filter-sterilized, neutralized 0.43 M cysteine·HCl·H<sub>2</sub>O as the reducing agent; 1.8 ml 0.5 M KHPO<sub>4</sub> buffer pH 6.9 and 2.67 ml 0.8 M NaHCO<sub>3</sub> as the carbon source, per litre diluted ASW. The thiosulfate, cysteine and NaHCO<sub>3</sub> were added after autoclaving and the pH adjusted to 7.1–7.2. The medium was dispensed under air as 10 ml aliquots into sterile 15×125 mm test tubes which were kept at 44 °C, inoculated, inverted several times and then quickly cooled on ice. Cells of S-1 grew as lens-shaped colonies at the OAI of this medium after 7–10 days, and several were aseptically extracted and serially diluted in shake tubes two more times.

After isolation, S-1<sup>T</sup> was routinely grown in a semi-solid growth medium containing (per l of ASW): 0.2 ml 0.2% aqueous resazurin, 10.0 ml 25% Na<sub>2</sub>S<sub>2</sub>O<sub>3</sub>·5H<sub>2</sub>O, 0.25 g NH<sub>4</sub>Cl, 2.0 ml neutralized 0.43 M cysteine·HCl·H<sub>2</sub>O, 2.67 ml 0.8 M NaHCO<sub>3</sub>, 5 ml modified Wolfe's mineral elixir [10 ], 0.5 ml vitamin solution [ 10 ] and 2.0 g Agar Noble (Difco Laboratories). The final pH was adjusted to 7.0–7.1. Cells grew as a microaerobic band of cells within the OAI below the surface of the medium ( Fig. 1a ). Although cysteine is a potential carbon and/or energy source, there was no observable growth of S-1<sup>T</sup> on the aforementioned medium lacking sulfide or thiosulfate, either with or without NaHCO<sub>3</sub>. Thus S-1<sup>T</sup> does not appear to be able to use cysteine as a carbon or energy source.

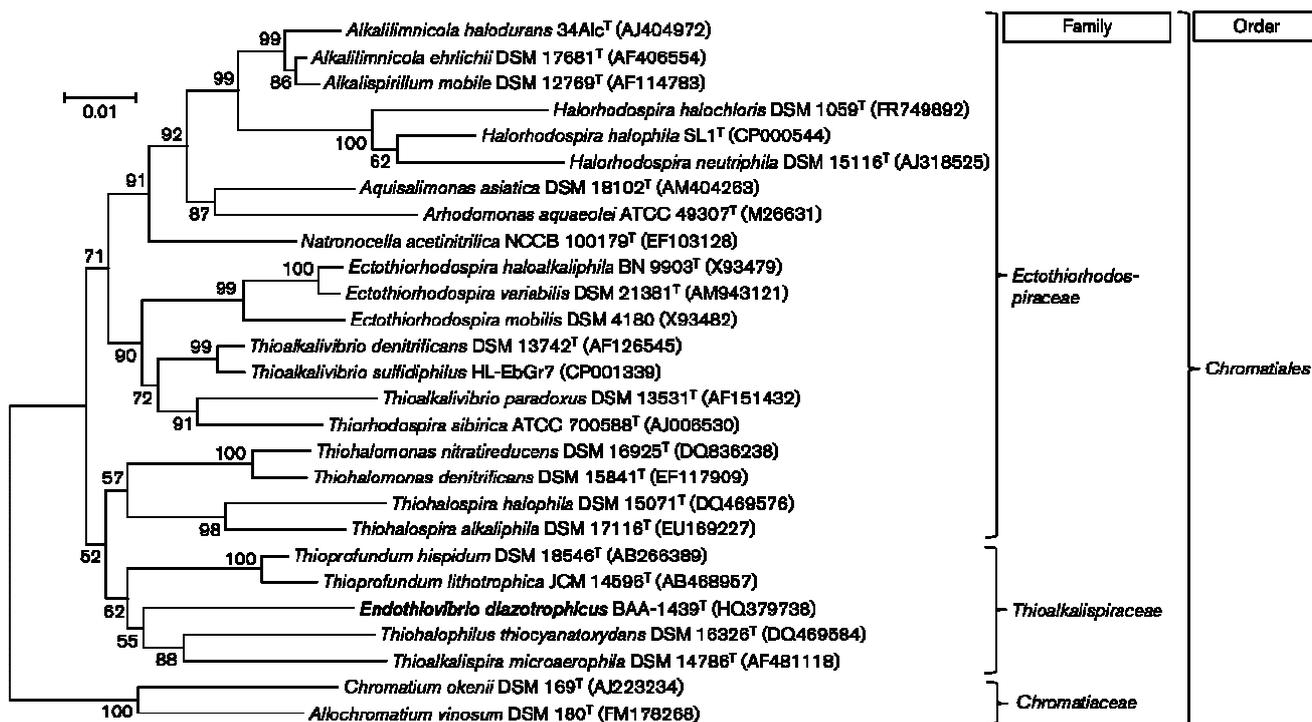


**Fig. 1.** (a) Growth of *S-1*<sup>T</sup> in semi-solid, thiosulfate, oxygen-gradient medium containing phenol red (two tubes at left) as a pH indicator or resazurin (two tubes at right) as an indicator of oxygen. Note that cells grow in inoculated tubes (2 and 4) as a microaerophilic band of cells below the meniscus. The color of tube 2 near the band of cells is yellow, indicating the production of acid, specifically sulfuric acid. In tube 4 cells grew within the oxic–anoxic transition zone as indicated by the resazurin which is colorless when reduced, in this case when oxygen is not present. The uninoculated tubes remained unchanged. (b) Differential interference contrast microscopic image of *S-1*<sup>T</sup> cells grown with thiosulfate as electron donor showing the presence of intracellular sulfur globules (c and d). Scanning transmission electron microscope (STEM) image of unstained cells of *S-1*<sup>T</sup> and corresponding O, P, S and Mg elemental maps of the same cells. One sulfur-rich and one phosphorus-rich inclusion is designated by labeled arrows in the STEM image. The map shows locations of numerous sulfur-rich globules and phosphorus-rich bodies that probably consist of polyphosphate. The lower cell appears to have lysed. Bars, 1  $\mu$ m.

To scale up liquid cultures for DNA extraction, *S-1*<sup>T</sup> was grown in liquid thiosulfate medium as described above, except that the volume of cysteine·HCl·H<sub>2</sub>O solution was decreased to 0.5 ml l<sup>-1</sup>, the concentration of NaHCO<sub>3</sub> was increased to 15 mM, and the medium was rendered anaerobic by sparging the medium with O<sub>2</sub>-free 7.5% CO<sub>2</sub> in N<sub>2</sub>. Oxygen was added to these bottles after autoclaving to a headspace concentration of 0.7% to initiate growth. Once turbidity (growth) became apparent O<sub>2</sub> was added in increments of 1% of the headspace when O<sub>2</sub> from the previous injection was utilized (when medium became colorless).

Determination of DNA G+C content was performed using HPLC by the Deutsche Sammlung von Mikroorganismen und Zellkulturen (DSMZ) according to the methods described by Cashion *et al.* [ 11 ], Tamaoka and Komagata [ 12 ], and Mesbah *et al.* [ 13 ]. The G+C content of genomic DNA of *S-1*<sup>T</sup>, as determined by HPLC, was 67.6±2.0 mol% ( *n*=6 replicates). The 16S rRNA gene sequence of *S-1*<sup>T</sup> was amplified from genomic DNA, using primers fD1 and rD1 according to the method of Weisburg *et al.* [ 14 ]. Using blastn, the 16S rRNA gene sequence of strain *S-1*<sup>T</sup> (GenBank accession number HQ379738) was shown to have the highest sequence identity to that of the rRNA gene from *Thioalkalivibrio sulfidiphilus* HL-EbGr7<sup>T</sup> (93% sequence identity; GenBank accession number CP001339), a member of the family *Ectothiorhodospiraceae*, among characterized strains. Phylogenetic analysis (Neighbor-Joining, based on 1078 aligned bp of 16S rRNA) recovered strain *S-1*<sup>T</sup> as nested within the family *Thioalkalispiraceae* of the

class *Gammaproteobacteria*, although *Thioalkalivibrio sulfidophilus* was found not to be the closest relative of S-1<sup>T</sup> (Fig. 2). This family contains obligate chemolithoautotrophs [6, 15–17], which accords with the chemolithoautotrophic metabolism of S-1<sup>T</sup>. S-1<sup>T</sup> was demonstrated only to be able to use sulfide and thiosulfate as electron donors for chemolithoautotrophic growth; other species of the family *Thioalkalispiraceae* have been shown to be able to use tetrathionate as an electron donor, including *Thiopfundum lithotrophicum* [17], *Thiopfundum hispidum* [6] and *Thioalkalispira microaerophila* [15], or to use thiocyanate, as in *Thiohalophilus thiocyanoxidans* [16]. S-1<sup>T</sup> only grew aerobically; other members of the *Thioalkalispiraceae* can respire using nitrate [6, 17] or nitrite [16], although (like S-1<sup>T</sup>) *Thioalkalispira microaerophila* only grows microaerobically [15]. As S-1<sup>T</sup> did not utilize any of a large number of organic compounds for growth, it appears to be an obligate chemolithoautotroph like other known members of the family *Thioalkalispiraceae* [6, 15–17].

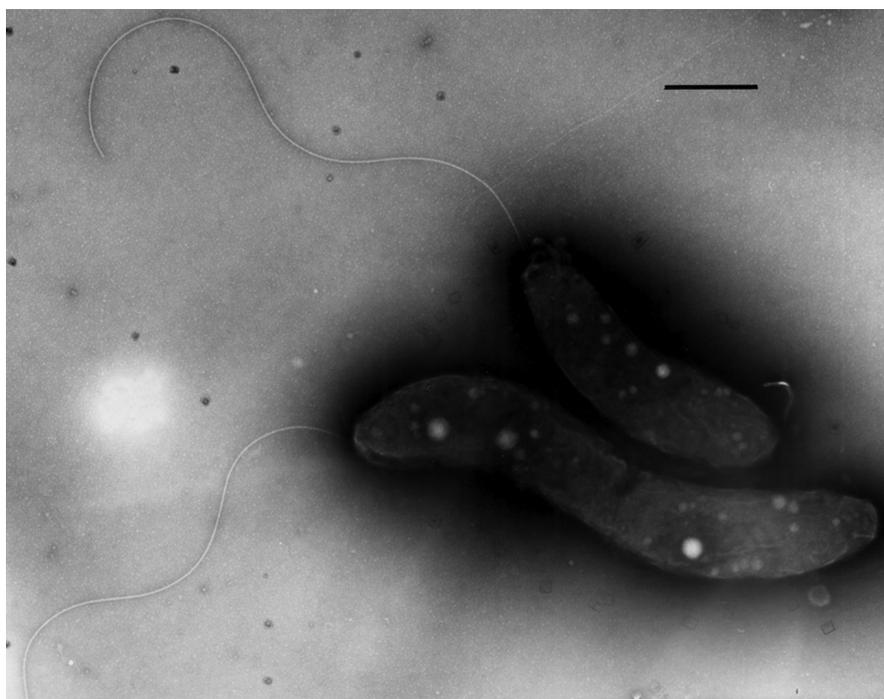


**Fig. 2.** Phylogenetic relationships of S-1<sup>T</sup> within the order *Chromatiales* of the class *Gammaproteobacteria* based on 16S rRNA gene sequences. S-1<sup>T</sup> was recovered within the family *Thioalkalispiraceae*. The phylogenetic tree was reconstructed using the Neighbor-Joining algorithm based on 1078 aligned bp of 16S rRNA gene sequences with sequences of *Chromatium okenii* and *Allochromatium vinosum* (family *Chromatiaceae*) used as outgroups. Bootstrap values (from 1000 replicates) higher than 50% are shown at nodes. Bar, 1 replacement per 100 nucleotide positions.

Analytical electron microscopy was performed on cells using a VG Microscopes model HB-5 scanning transmission electron microscope (STEM; Fisons Instrument Surface Systems) operating at 100 kV linked to a field-emission electron gun, a Link LZ-5 X-ray detector (Link Analytical) and an AN10000 X-ray analysis system. Motility of S-1<sup>T</sup> was determined using a Zeiss AxioImager M1 light microscope (Carl Zeiss MicroImaging) equipped with phase-contrast and differential interference contrast capabilities.

Cells of S-1<sup>T</sup> are Gram-stain-negative rods that are straight to curved in morphology ( Figs 1b, 3 ). Cells have an average length of 4.9±1.1 µm and a width of 1.2±0.2 µm ( n=57). Cells are motile by means of a single, polar, unsheathed flagellum ( Fig. 3 ). The average swimming speed for cells of S-1<sup>T</sup> in fresh culture was 40.3±8.2 µm s<sup>-1</sup> ( n=61). Cells grew chemolithoautotrophically on sulfide and thiosulfate as electron donors, and produced large internal sulfur globules and phosphate-rich

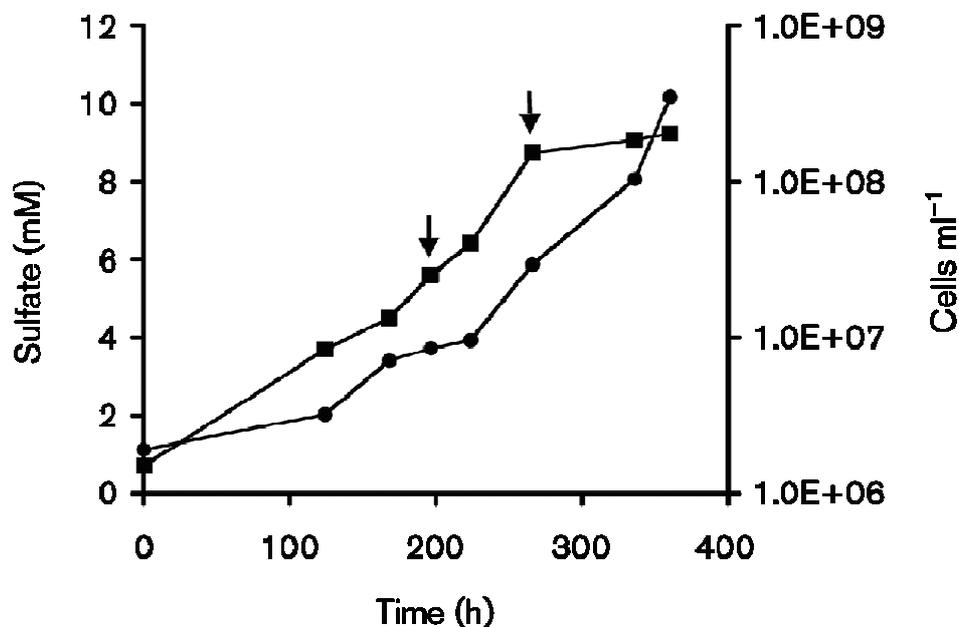
deposits, presumably consisting of polyphosphate ( Fig. 1b–d ). These intracellular bodies were observed to be larger when cells were grown on thiosulfate than when they were grown on sulfide (data not shown). Such internal sulfur deposits are observed in some other members of the family *Thioalkalispiraceae* , and their presence is variable within this family [ 6, 18 ].



**Fig. 3.** *Transmission electron microscope (TEM) image of cells of S-1<sup>T</sup> grown with sulfide as the electron donor, negatively-stained with 0.5% uranyl acetate. Note the presence of a single, polar, unsheathed flagellum. Bar, 1 μm.*

During growth of S-1<sup>T</sup> in thiosulfate medium, internal sulfur globules disappeared as thiosulfate became depleted in the growth medium and the pH dropped to about 5.4.

Sulfate concentrations were measured turbidometrically using the barium gel method of Tabatabai [ 19 ]. Sulfate was released by cells of S-1<sup>T</sup> only during growth on thiosulfate ( Fig. 4 ). The evidence is consistent with elemental sulfur storage being an obligate intermediate in the oxidation of sulfide and thiosulfate by cells of S-1<sup>T</sup> during thioautotrophic growth. The release of sulfate into the medium is consistent with the sulfane group of thiosulfate being transformed into storage sulfur, whereas the sulfone group is immediately converted to sulfate and excreted [ 2, 20, 21 ]. As exogenous thiosulfate appeared to be consumed by S-1<sup>T</sup> cultures, the cells utilized internal stored sulfur, which accordingly resulted in the depletion of intracellular sulfur globules. Moreover, when resazurin was replaced with phenol red in thiosulfate-containing growth medium, the medium turned from pink to yellow, particularly around the band of cells, indicating that the strains oxidize thiosulfate to sulfuric acid (sulfate; Fig. 1a ).



**Fig. 4.** Growth (■) and sulfate production (□) from thiosulfate in a liquid culture of *S-1<sup>T</sup>* grown microaerobically. Initial  $O_2$  concentration 1.3% of the headspace (20 ml) and increments of 2% of the headspace was added to the culture at times indicated by the arrows. Sulfate was never formed in the absence of thiosulfate.

*S-1<sup>T</sup>* appears to be well adapted to the marine environment and grew in 10–100% ASW using the ASW composition of Lyman and Fleming [22]. Sodium chloride alone did not substitute for the marine salts present in ASW in concentrations of 1.7, 2.6 and 3.5% (Table 1). Cells grew in media with a pH range of 5.2–8.0 with an apparent optimum growth pH of about 7.2 (the pH of the sample from which it was isolated) (Table 1). Catalase activity was determined by generation of  $O_2$  gas bubbles when cell suspensions were added to aqueous 3%  $H_2O_2$  solution. Oxidase activity was tested with the *N, N, N', N'*-tetramethyl-*p*-phenylenediamine dihydrochloride assay using oxidase strips (Sigma-Aldrich). *S-1<sup>T</sup>* was shown to be catalase-negative and oxidase-negative. For carbon and energy source testing, thiosulfate (as  $Na_2S_2O_3 \cdot 5H_2O$ ) was omitted from the semi-solid growth medium and a carbon source was substituted at a concentration of 0.1% (w/v or v/v; sodium salts used for all acids, neutralized when necessary; l-enantiomers were used for amino acids, d-enantiomers for sugars). The following were tested as carbon sources for *S-1<sup>T</sup>*: citrate, 2-oxoglutarate, malate, pyruvate, succinate, acetate, butyrate, glycolate, glyoxylate, lactate, malonate, propionate, quinate, salicylate, tartrate, valerate, urea, alanine, sodium glutamate, isoleucine, leucine, lycine, methionine, phenylalanine, proline, serine, threonine, tryptophan, valine, butanol, ethanol, glycerol, isopropanol, methanol, acetaldehyde, formaldehyde, adonitol (ribitol), aesculin, amygdalin, arabinose, cellobiose, dulcitol, fructose, fucose, galactose, gluconate, glucose, glycogen, hippurate, inulin, lactose, lyxose, maltose, mannitol, mannose, melezitose, melibiose, raffinose, rhamnose, ribose, salicin, sedoheptulose, sorbitol, sorbose, sucrose, trehalose, xylose, casamino acids, peptone, tryptone, yeast extract and gelatin. Cells for inoculation into carbon testing medium were aseptically centrifuged at 10000 *g* at 4 °C for 20 min and washed once with sterile 20 mM Tris·HCl in ASW, recentrifuged and resuspended in the same buffer to a cell concentration of  $1.0 \times 10^8$  cells ml<sup>-1</sup>. Each tube containing 10 ml medium was inoculated with 0.1 ml of this cell suspension (final cell concentration =  $1.0 \times 10^6$  cells ml<sup>-1</sup>) which was distributed evenly throughout the tube by inverting the tubes several times at 44 °C prior to placing the tubes quickly in an ice bath for solidification. Anaerobic growth was determined similarly except that cultures were sparged with  $O_2$ -free  $N_2$ . All cultures were incubated at 28 °C. Growth was determined by measuring cell numbers as direct cell counts. Tubes containing semi-solid agar where cells grew as a microaerophilic band of cells at the OAI were inverted quickly at least 20 times to evenly disperse the cells before a sample was removed for counting cells as described above. Cells of *S-1<sup>T</sup>* did not grow with any of the carbon sources

tested and thus we infer that S-1<sup>T</sup> is either not capable of chemoorganoheterotrophic growth or has an extremely limited capability of doing so.

Characteristic	S-1 <sup>T</sup>	<i>Thioalkalispira microaerophila</i> ALEN 1 <sup>T</sup>	<i>Thiohalophilus thiocyanoxidans</i> HRhD 2 <sup>T</sup>	<i>Thiopfundum lithotrophicum</i> 106 <sup>T</sup>	<i>Thiopfundum hispidum</i> gps61 <sup>T</sup>
Cell morphology	Rod or vibrioid	Spiral rod	Rod	Spiral rod	Rod
Motility	+	+	-	+	-
	(Single polar flagellum)	(Single polar flagellum)		(Single polar flagellum)	
Intracellular sulfur deposits	+	-	ND	ND	-
O <sub>2</sub> requirement	Microaerobic	Microaerobic	Facultatively anaerobic	Facultatively anaerobic	Facultatively anaerobic
Terminal electron acceptors	O <sub>2</sub>	O <sub>2</sub>	O <sub>2</sub> , NO <sub>2</sub>	O <sub>2</sub> , NO <sub>3</sub>	O <sub>2</sub> , NO <sub>3</sub>
Electron donors	S <sup>2-</sup> , S <sub>2</sub> O <sub>3</sub> <sup>2-</sup>	S <sup>2-</sup> , S <sub>8</sub> <sup>2-</sup> , S <sup>0</sup> , S <sub>2</sub> O <sub>3</sub> <sup>2-</sup>	S <sup>2-</sup> , S <sub>2</sub> O <sub>3</sub> <sup>2-</sup> , SCN <sup>-</sup>	S <sup>0</sup> , S <sub>2</sub> O <sub>3</sub> <sup>2-</sup> , S <sub>4</sub> O <sub>6</sub> <sup>2-</sup> , SO <sub>3</sub> <sup>2-</sup> ‡	S <sup>0</sup> , S <sub>2</sub> O <sub>3</sub> <sup>2-</sup> , S <sub>4</sub> O <sub>6</sub> <sup>2-</sup>
Metabolism	Obligate chemolithoautotroph	Obligate chemolithoautotroph	Obligate chemolithoautotroph	Obligate chemolithoautotroph	Obligate chemolithoautotroph
Phototrophy	-	-	ND	ND	-
N <sub>2</sub> fixation	+	ND	ND	+	ND
Oxidase	-	+	ND	ND	+
Catalase	-	+	ND	ND	-
Growth temperature range (°C)	4-37.5 (optimum 28)	30†	30†	30-55 (optimum 50)	29-43 (optimum 39)
Growth pH range	5.2-8.0 (-7.2)	8.0-10.4 (optimum ~10)	6.5-8.2 (optimum 7.5)	5.8-7.6 (optimum 7.0)	6.0-8.0 (optimum 7.0)
Growth NaCl range (mol l <sup>-1</sup> )	*	0.2-1.4 (optimum 0.5)	1.0-4.0 (optimum 1.5)	0.21-0.72 (optimum 0.51)§	0.17-0.68 (optimum 0.34)§
DNA G+C content (mol %)	67.6	58.9	58.2	65.7	62.9

\*Cells did not grow when NaCl was the only salt present in cultures at 0.29, 0.44 and 0.60 mol l<sup>-1</sup>. They did grow, however, in concentrations of an artificial seawater (composition from [22]) from 10 to 100 % but not below 10 %.  
 †Growth temperature only, not optimum temperature or temperature range, was reported. ‡*Thiopfundum lithotrophicum* 106T was able to use sulfite as an electron donor in the analysis of Takai et al. [17], but was not able to use sulfite as an electron donor in the analysis of Mori et al. [6]. §Concentration values originally reported as % (w.v), and converted here to mol l<sup>-1</sup>

**Table 1.** Characteristics that differentiate *Endothiovibrio diazotrophicus* S-1<sup>T</sup> from other members of the family *Thioalkalispiraceae*.

Reference data are from [15, 16], Takai et al. [17] and Mori et al. [6]. +, positive; -, negative; nd, not determined.

Anaerobic growth (using thiosulfate as the electron donor) was tested in semi-solid medium using the following as terminal electron acceptors (sodium salts, where appropriate): NO<sub>3</sub><sup>-</sup> (5 and 10 mM); NO<sub>2</sub><sup>-</sup> (2 mM); N<sub>2</sub>O (1 atm); fumarate (20 mM); SO<sub>4</sub><sup>2-</sup> (5 and 10 mM); trimethylamine oxide (15 mM); and dimethylsulfoxide (15 mM). Cells only grew when oxygen was present and did not grow anaerobically using any of the tested electron acceptors aside from oxygen. Consequently, it appears that S-1<sup>T</sup> is an obligate aerobe.

Alternative electron donors for S-1<sup>T</sup> were tested microaerobically using semi-solid medium as described above, except that elemental sulfur (5 and 10 mM), tetrathionate (10 mM) and thiocyanate (10 mM) were used instead of thiosulfate. No growth was observed on any of these compounds. Thus, sulfide and thiosulfate are the only compounds that we could demonstrate supported growth of S-1<sup>T</sup>.

The contribution of CO<sub>2</sub> to total cellular carbon was measured as <sup>14</sup>C-bicarbonate incorporation into cell material. Briefly, sterile, molten semi-solid medium containing 10 mM thiosulfate as the electron source was supplemented with <sup>14</sup>C-bicarbonate (typically 1.0 mCi ml<sup>-1</sup>, averaging 50 mCi mmol<sup>-1</sup>; New England Nuclear). Samples were removed for measurement of: (1) total counts per ml and (2) total inorganic carbon which was determined by acidification of an aliquot of growth medium and measuring the CO<sub>2</sub> evolved using a LIRA infrared gas analyzer. The final specific activity in the

growth medium was typically between  $4.5$  and  $6.0 \times 10^4$  counts  $\text{min}^{-1} \mu\text{g}$  inorganic  $\text{C}^{-1}$ . Treatment and fractionation of cells, protein determinations, controls and kinetic isotope correction were as previously described by Nelson and Jannasch [ 9 ]. Liquid scintillation counting was accomplished using a model LS 6500 (Beckman Instruments) liquid scintillation counter. S-1<sup>T</sup> cells clearly incorporated  $^{14}\text{C}$  from  $\text{H}^{14}\text{CO}_3^- / ^{14}\text{CO}_2$  in growth experiments in which radiolabelled  $\text{H}^{14}\text{CO}_3^- / ^{14}\text{CO}_2$  was the sole carbon source (Table S1, available in the online Supplementary Material). However, because heterotrophs assimilate  $\text{CO}_2$  anaerobically, to prove autotrophy in an organism it must be shown that most if not all the cellular carbon is derived from  $\text{CO}_2$ . To do this, we measured the amount of protein in the separate experimental cultures and then calculated the amount of protein carbon using an average C content of protein of 54% [ 23, 24 ]. For each experiment, the amount of  $^{14}\text{C}$  from  $\text{H}^{14}\text{CO}_3^- / ^{14}\text{CO}_2$  incorporated into protein was compared with the calculated amount of protein carbon (Table S1). The results indicated that the average percentage of protein carbon derived from  $\text{H}^{14}\text{CO}_3^- / ^{14}\text{CO}_2$  was  $97.6 \pm 3.2\%$  for S-1<sup>T</sup>. Our results indicate that essentially all cellular carbon in S-1<sup>T</sup> comes from  $\text{H}^{14}\text{CO}_3^- / ^{14}\text{CO}_2$ , a result consistent with autotrophic growth.

Ribulose-1,5-bisphosphate carboxylase/oxygenase (RubisCO) activity of cell-free extracts was determined as described by Beudeker *et al.* [ 25 ] except that the dithiothreitol concentration was changed to 5 mM and the pH adjusted to either 7.2 or 8.2. Cell-free extracts were prepared by harvesting cells by centrifugation as described above, suspending cells in 20 mM Tris HCl, pH 7.2 or 8.2, and passing this suspension through a French pressure cell. Supernatant was obtained from the crude extracts by centrifugation at 10000 *g* at 4 °C for 20 min. Cell-free extracts from S-1<sup>T</sup> grown microaerobically under autotrophic conditions with thiosulfate as the electron donor and  $\text{HCO}_3^-$  as the sole C source showed RubisCO activity ( $9.6 \pm 2.3$  and  $5.4 \pm 2.1$  nmol  $\text{CO}_2$  fixed  $\text{min}^{-1} \text{mg}$  protein<sup>-1</sup> at pH 7.2 and 8.2, respectively; (Table S2). Fixation of  $\text{CO}_2$  in the assays was always ribulose bisphosphate (substrate)-dependent (data not shown).

The partial sequences of the RubisCO forms I ‘green’ (*cbbL*) and II (*cbbM*) genes were amplified using the degenerate primer pairs RubIgF–RubIgR and RuIIF1–RuIIF3 [ 26 ], respectively. Primer synthesis and DNA sequencing was performed at Iowa State University’s DNA Synthesis and Sequencing Facility. Sequences of genes for the RubisCO subunits CbbL and CbbM were obtained (GenBank accession numbers JN676153 and JN676154), which confirms that S-1<sup>T</sup> employs the Calvin–Benson–Bassham cycle for autotrophic growth.

For the determination of  $^{15}\text{N}_2$  fixation, cells were grown microaerobically in 155 ml serum vials containing 55 ml nitrogen-free growth medium. The headspace gas was Ar with 1%  $\text{O}_2$ . Additions to the growth medium were 4 mM  $\text{NH}_4\text{Cl}$  and/or 10 ml  $^{14}\text{N}_2$  or 10 ml  $^{15}\text{N}_2$  (99 atomic %  $^{15}\text{N}$ ; Isotec). Cells were harvested by centrifugation (10000 *g*, 4 °C, 20 min) after turbidity became apparent, then washed twice in cold 20 mM Tris buffer in diluted artificial sea water (pH 7.0), and freeze dried. The atomic percentage  $^{15}\text{N}$  of freeze dried cells was determined using a Model NA-1500 elemental analyzer (Fisons Instruments) linked to a Finnegan MAT Delta S isotope ratio mass spectrometer (ThermoQuest). Nitrogenase activity of whole cells was determined as acetylene ( $\text{C}_2\text{H}_2$ ) reduction to ethylene ( $\text{C}_2\text{H}_4$ ) in  $\text{O}_2/\text{S}_2\text{O}_3^{2-}$  semi-solid gradient cultures as previously described [ 27 ]. Cells of S-1<sup>T</sup> fixed  $^{15}\text{N}_2$  when a fixed nitrogen source was absent from the growth medium;  $^{15}\text{N}_2$  fixation was inhibited when 4 mM  $\text{NH}_4\text{Cl}$  was included in the growth medium (Table S3). Cells of S-1<sup>T</sup> also showed relatively high nitrogenase activities (Table S4) as measured by acetylene ( $\text{C}_2\text{H}_2$ ) reduction to  $\text{C}_2\text{H}_4$  in  $\text{O}_2/\text{S}_2\text{O}_3^{2-}$  semi-solid gradient cultures.  $\text{C}_2\text{H}_2$  reduction was inhibited by the addition of 4 mM  $\text{NH}_4\text{Cl}$  but not by 4 mM  $\text{NaNO}_3$ .

Fatty acid analyses were carried out by the Identification Service of the DSMZ GmbH (Braunschweig, Germany). Fatty acids were extracted, purified, methylated and quantified by gas chromatography using the standard Microbial Identification System [ 28, 29 ]. Liquid cultures grown on thiosulfate under microaerobic conditions were used, with the same composition as ASW-based semi-solid medium (see above), except that agar was withheld. Although it is difficult to absolutely

determine what stage of growth cells of S-1<sup>T</sup> are in when grown in liquid, based on the observed growth, cells for these analyses were harvested about 24 h after a large injection of O<sub>2</sub> into the culture and the culture was therefore probably in late exponential phase. The major cellular fatty acids were C<sub>16:1</sub>ω7 c (37% of the total fatty acids), C<sub>16:0</sub> (30%), and C<sub>18:1</sub>ω7 c (22%). The high proportions of the fatty acids C<sub>16:1</sub>ω7 and C<sub>16:0</sub> are consistent with the cellular fatty acid composition of other species of the family *Thioalkalispiraceae* [ 6, 16, 17 ], with the caveat that for the piezophilic species *Thiopfundum lithotrophicum* the unsaturated fatty acid C<sub>16:1</sub>ω7 exhibited an increased proportion of total fatty acids with increased hydrostatic pressure [ 17 ]. S-1<sup>T</sup> also contained C<sub>18:0</sub> (5.0%), C<sub>18:2</sub>ω6,9 c (1.3%), C<sub>18:1</sub>ω5 c (1.2%), C<sub>17:0</sub> (1.0%) and C<sub>16:1</sub>ω5 c (0.5%), C<sub>14:0</sub> (0.4%), C<sub>17:1</sub>ω6 c (0.4%), and C<sub>15:0</sub> (0.3%) as minor fatty acid components.

Analysis of respiratory quinones and polar lipids were carried out by the Identification Service and Dr Brian Tindall, DSMZ (Braunschweig, Germany). Cells for these analyses were grown in liquid culture as described earlier. Q8 was found to be the only respiratory quinone. Major polar lipids were phosphatidylethanolamine and phosphatidylglycerol, as well as an unknown phospholipid (Fig. S1).

#### DESCRIPTION OF ENDOTHIOVIBRIO GEN. NOV.

*Endothiovibrio* [En.do.thi.o.vi'bri.o. Gr. pref. *endo* within; Gr. n. *theion* (Latin transliteration *thium*) sulfur; N.L. masc. n. *Vibrio* a bacterial genus; N.L. masc. n. *Endothiovibrio* vibrio with internal sulfur].

Members of this genus belong to the family *Thioalkalispiraceae* within the class *Gammaproteobacteria*. Cells are Gram-stain-negative rods that are straight to curved in morphology and motile by means of a single polar unsheathed flagellum. The Calvin–Benson–Bassham pathway is used for CO<sub>2</sub> fixation. The type species is *Endothiovibrio diazotrophicus*.

#### DESCRIPTION OF ENDOTHIOVIBRIO DIAZOTROPHICUS SP. NOV.

*Endothiovibrio diazotrophicus* (di.a.zo.tro'phi.cus. Gr. pref. *di* two, double; N. L. n. *azotum* nitrogen; Gr. adj. *trophikos* tending or feeding; N. L. masc. adj. *diazotrophicus*, one that feeds on dinitrogen).

Description as for genus, with the following additional characteristics. Cells have an average length of 4.9±1.1 μm and a width of 1.2±0.2 μm. Doubling time approximately 27 h. Cells grow chemolithoautotrophically, possibly obligately, on sulfide and thiosulfate as the electron donors, only under microaerobic conditions. When grown on thiosulfate or sulfide, cells produced large internal sulfur globules and phosphate-rich (putative polyphosphate) deposits. Diazotrophic, with demonstrable nitrogenase activity. Nitrogen fixation is completely inhibited by NH<sub>4</sub>Cl. Mesophilic, with a growth temperature range of 5–37 °C, and optimal growth occurring at around 28 °C. Can grow in medium containing 10–100% artificial seawater, but not below 10%, and NaCl alone cannot substitute for sea salts. Major cellular fatty acids are C<sub>16:1</sub>ω7c, C<sub>16:0</sub>, and C<sub>18:1</sub>ω7c. Major respiratory quinone is Q8. Phosphatidylethanolamine and phosphatidylglycerol are the dominant polar lipids. Catalase-negative. Oxidase-negative.

The type strain is S-1<sup>T</sup> (=ATCC BAA-1439<sup>T</sup>=JCM 17961<sup>T</sup>), originally isolated from School Street Marsh at Woods Hole, Massachusetts, USA. The DNA G + C content of the type strain is 67.6±2.0 mol%.

TheGenBank accession numbers for the 16S rRNA, *cbbL* and *cbbM* gene sequences of *Endothiovibriodiazotrophicus* S-1<sup>T</sup> are HQ379738, JN676153, and JN676154, respectively.

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## ABBREVIATIONS

ASW, artificial seawater; OAI, oxic–anoxic interface; RubisCO, ribulose biphosphate 30 carboxylase/oxygenase; STEM, scanning transmission electron microscope; TEM, transmission electron microscope.

## REFERENCES

1. Garrity GM, Bell JA, Lilburn TG. Class III. *Gammaproteobacteria* class. nov. In Brenner DJ, Krieg NR, Staley JT, Garrity GM (editors) *Bergey's Manual of Systematic Bacteriology (2nd ed)* 2nd ed. vol. 2 New York: Springer; 2005; pp.:1–850
2. Dahl C, Prange A. Bacterial sulfur globules: occurrence, structure and metabolism. In Shively JM (editor) *Inclusions in Prokaryotes* Berlin: Springer Verlag; 2006 pp.:21–51.
3. Frigaard NU, Dahl C. Sulfur metabolism in phototrophic sulfur bacteria. *Adv Microb Physiol* 2009;54:103–200. doi:[http://dx.doi.org.inee.bib.cnrs.fr/10.1016/S0065-2911\(08\)00002-7](http://dx.doi.org.inee.bib.cnrs.fr/10.1016/S0065-2911(08)00002-7).
4. Pott AS, Dahl C. Sirohaem sulfite reductase and other proteins encoded by genes at the *dsr* locus of *Chromatium vinosum* are involved in the oxidation of intracellular sulfur. *Microbiol* 1998;144:1881–1894. doi: <http://dx.doi.org.inee.bib.cnrs.fr/10.1099/00221287-144-7-1881>.
5. Lee K, Lee HK, Choi TH, Kim KM, Cho JC. *Granulosicoccaceae* fam. nov., to include *Granulosicoccus antarcticus* gen. nov., sp. nov., a non-phototrophic, obligately aerobic chemoheterotroph in the order *Chromatiales*, isolated from Antarctic seawater. *J Microbiol Biotechnol* 2007;17:1483–1490.
6. Mori K, Suzuki K, Urabe T, Sugihara M, Tanaka K, et al. *Thiopfundum hispidum* sp. nov., an obligately chemolithoautotrophic sulfur-oxidizing gammaproteobacterium isolated from the hydrothermal field on Suiyo Seamount, and proposal of *Thioalkalispiraceae* fam. nov. in the order *Chromatiales*. *Int J Syst Evol Microbiol* 2011;61:2412–2418. doi: <http://dx.doi.org.inee.bib.cnrs.fr/10.1099/ijms.0.026963-0>.
7. Wang G, Tang M, Li T, Dai S, Wu H, et al. *Wenzhouxiangella marina* gen. nov, sp. nov, a marine bacterium from the culture broth of *Picochlorum* sp. 122, and proposal of *Wenzhouxiangellaceae* fam. nov. in the order *Chromatiales*. *Antonie Van Leeuwenhoek* 2015;107:1625–1632. doi: <http://dx.doi.org.inee.bib.cnrs.fr/10.1007/s10482-015-0458-7>.
8. Du ZJ, Wang ZJ, Zhao J X, Chen GJ. *Woeseia oceani* gen. nov., sp. nov., a chemoheterotrophic member of the order *Chromatiales*, and proposal of *Woeseiaceae* fam. nov. *Int J Syst Evol Microbiol* 2016;66:107112. doi:<http://dx.doi.org.inee.bib.cnrs.fr/10.1099/ijsem.0.000683>.
9. Nelson DC, Jannasch HW. Chemoautotrophic growth of a marine *Beggiatoa* in sulfide-gradient cultures. *Arch Microbiol* 1983; 136: 262–269. doi: <http://dx.doi.org.inee.bib.cnrs.fr/10.1007/BF00425214>.
10. Frankel RB, Bazylinski DA, Johnson MS, Taylor BL. Magneto-aerotaxis in marine coccoid bacteria. *Biophys J* 1997;73:994–1000. doi:[http://dx.doi.org.inee.bib.cnrs.fr/10.1016/S0006-3495\(97\)78132-3](http://dx.doi.org.inee.bib.cnrs.fr/10.1016/S0006-3495(97)78132-3).
11. Cashion P, Holder-Franklin MA, McCully J, Franklin M. A rapid method for the base ratio determination of bacterial DNA. *Anal Biochem* 1977; 81: 461–466. doi: [http://dx.doi.org.inee.bib.cnrs.fr/10.1016/0003-2697\(77\)90720-5](http://dx.doi.org.inee.bib.cnrs.fr/10.1016/0003-2697(77)90720-5).

12. Tamaoka J, Komagata K. Determination of base composition by reversed-phase high-performance liquid chromatography. *FEMS Microbiol Lett* 1984;25:125–128. doi: <http://dx.doi.org.inee.bib.cnrs.fr/10.1111/j.1574-6968.1984.tb01388.x>.
13. Mesbah M, Premachandran U, Whitman W. Precise measurement of the G+C content of deoxyribonucleic acid by high performance liquid chromatography. *Int J Syst Bacteriol* 1989;39:159–167. doi: <http://dx.doi.org.inee.bib.cnrs.fr/10.1099/00207713-39-2-159>.
14. Weisburg WG, Barns SM, Lane DJ. 16S ribosomal DNA amplification for phylogenetic study. *J Bacteriol* 1991;173:697–703. doi:<http://dx.doi.org.inee.bib.cnrs.fr/10.1128/jb.173.2.697-703.1991..>
15. Sorokin DY, Tourova TP, Kolganova T V, Sjollema KA, Kuenen JG. *Thioalkalispira microaerophila* gen. nov., sp. nov., a novel lithoautotrophic, sulfur-oxidizing bacterium from a soda lake. *Int J Syst Evol Microbiol* 2002;52:2175–2182. doi:<http://dx.doi.org.inee.bib.cnrs.fr/10.1099/00207713-52-6-2175>.
16. Sorokin DY, Tourova TP, Bezsoudnova EY, Pol A, Muyzer G. Denitrification in a binary culture and thiocyanate metabolism in *Thiohalophilus thiocyanoxidans* gen. nov. sp. nov. – a moderately halophilic chemolithoautotrophic sulfur oxidizing gammaproteobacterium from hypersaline lakes. *Arch Microbiol* 2007;187:441–450. doi: <http://dx.doi.org.inee.bib.cnrs.fr/10.1007/s00203-006-0208-3>.
17. Takai K, Miyazaki M, Hirayama H, Nakagawa S, Querellou J, *et al.* Isolation and physiological characterization of two novel, piezophilic, thermophilic chemolithoautotrophs from a deep-sea hydrothermal vent chimney. *Environ Microbiol* 2009;11:1983–1997. doi:<http://dx.doi.org.inee.bib.cnrs.fr/10.1111/j.1462-2920.2009.01921.x>.
18. Sorokin DY, Lysenko AM, Mityushina LL, Tourova TP, Jones BE, *et al.* *Thioalkalimicrobium aerophilum* gen. nov., sp. nov. and *Thioalkalimicrobium sibericum* sp. nov., and *Thioalkalivibrio versutus* gen. nov., sp. nov., *Thioalkalivibrio nitratis* sp. nov., novel and *Thioalkalivibrio denitrificans* sp. nov., novel obligately alkaliphilic and obligately chemolithoautotrophic sulfur-oxidizing bacteria from soda lakes. *Int J Syst Evol Microbiol* 2001;51:565–580. doi: <http://dx.doi.org.inee.bib.cnrs.fr/10.1099/00207713-51-2-565>.
19. Tabatabai MA. A rapid method for determination of sulfate in water samples. *Environ Lett* 1974;7:237–243. doi:<http://dx.doi.org.inee.bib.cnrs.fr/10.1080/00139307409437403>.
20. Hensen D, Sperling D, Trüper HG, Brune DC, Dahl C. Thiosulfate oxidation in the phototrophic sulfur bacterium *Allochromatium vinosum*. *Mol Microbiol* 2006;62:794–810. doi: <http://dx.doi.org.inee.bib.cnrs.fr/10.1111/j.1365-2958.2006.05408.x..>
21. Franz B, Gehrke T, Lichtenberg H, Hormes J, Dahl C, *et al.* Unexpected extracellular and intracellular sulfur species during growth of *Allochromatium vinosum* with reduced sulfur compounds. *Microbiol* 2009;155:2766–2774. doi:<http://dx.doi.org.inee.bib.cnrs.fr/10.1099/mic.0.027904-0>.
22. Lyman J, Fleming RH. Composition of seawater. *J Mar Res* 1940;3:134–146.
23. Jukes TH, Homquist R, Moise H. Amino acid composition of proteins: selection against the genetic code. *Science* 1975;189:50–51. doi:<http://dx.doi.org.inee.bib.cnrs.fr/10.1126/science.237322>.
24. Doolittle RF. Similar amino acid sequences: chance or common ancestry? *Science* 1981;214:149\_159. doi:<http://dx.doi.org.inee.bib.cnrs.fr/10.1126/science.7280687>.
25. Beudeker RF, Cannon GC, Kuenen JG, Shively JM. Relations between d-ribulose-1,5-bisphosphate carboxylase, carboxysomes and CO<sub>2</sub> fixing capacity in the obligate chemolithotroph *Thiobacillus neapolitanus* grown under different limitations in the chemostat. *ArchMicrobiol* 1980;124:185\_189. doi: <http://dx.doi.org.inee.bib.cnrs.fr/10.1007/BF00427725>.

26. Spiridonova EM, Berg IA, Kolganova TV, Ivanovsky RN, Kuznetsov BB, *et al.* An Oligonucleotide primer System for amplification of the ribulose-1,5-bisphosphate carboxylase/oxygenase genes of bacteria of various Taxonomic groups. *Microbiol-USSR* 2004;73:316\_325. doi:<http://dx.doi.org.inee.bib.cnrs.fr/10.1023/B:MICL.0000032243.93917.30>.
27. Bazyliniski DA, Dean AJ, Schüler D, Phillips EJP, Lovley DR. N<sub>2</sub>-dependent growth and nitrogenase activity in the metal-metabolizing bacteria, *Geobacter* and *Magnetospirillum* species. *Environ Microbiol* 2000;2:266–273. doi:<http://dx.doi.org.inee.bib.cnrs.fr/10.1046/j.1462-2920.2000.00096.x>.
28. Sasser M. *Identification of Bacteria by Gas Chromatography of Cellular Fatty Acids (MIDI Technical Note 101)*. MIDI Technical Note 101. Newark, DE: MIDI Inc; 1990.
29. Kämpfer P, Kroppenstedt RM. Numerical analysis of fatty acid patterns of coryneform bacteria and related taxa. *Can J Microbiol* 1996;42:989–1005. doi: <http://dx.doi.org.inee.bib.cnrs.fr/10.1139/m96-128>.