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# Characterization of Desulfovibrio fructosovorans sp. nov.

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Abstract. Desulfovibrio strain JJ isolated from estuarine sediment differed from all other described Desulfovibrio species by the ability to degrade fructose. The oxidation was incomplete, leading to acetate production. Fructose, malate and fumarate were fermented mainly to succinate and acetate in the absence of an external electron acceptor. The pH and temperature optima for growth were 7.0 and 35°C respectively. Strain JJ was motile by means of a single polar flagellum. The DNA base composition was 64.13% G+C. Cytochrome  $c_3$  and desulfoviridin were present. These characteristics established the isolate as a new species of the genus Desulfovibrio, and the name Desulfovibrio fructosovorans is proposed.

**Key words:** Sulfate reduction – Fermentation – Fructose – *Desulfovibrio fructosovorans* sp. nov.

Among the sulfate-reducing bacteria, the genus Desulfovibrio was known to have a very limited range of oxidizable substrates including hydrogen, ethanol, lactate, formate, malate, fumarate and succinate (Postgate 1979). Recently, Desulfovibrio strains were isolated that utilize amino acids (Stams et al. 1985), methanol and other alcohols (Braun and Stolp 1985; Nanninga and Gottschal 1986). The newly described Desulfovibrio sapovorans and D. baarsii differed from the formerly known "classical" Desulfovibrio species by the fact they used fatty acids (Widdel 1980; Widdel and Pfennig 1984).

There are few reports on the utilization of sugars by sulfate-reducers. *Desulfotomaculum nigrificans* is the only sulfate-reducing bacterium described so far that uses fructose whether sulfate is present or not (Klemps et al. 1985). In the present paper, we report on a new strain of the "classical"  $H_2$ -oxidizing *Desulfovibrio* group.

# Materials and methods

# Source of organism

*Desulfovibrio* strain JJ was kindly provided by W. J. Jones, University of Illinois at Urbana Champaign, USA. It was isolated from estuarine sediment and was used as hydrogen oxidizing bacterium in a defined mixed bacterial culture

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on sucrose (Jones et al. 1984). It was tentatively named D. vulgaris strain JJ. Further experiments showed that strain JJ metabolized fructose (Cord-Ruwisch et al. 1986). The fructose grown culture has been serially diluted and inoculated into roll tubes containing  $H_2$ , fructose or lactate. Picked colonies were able to grow on fructose (Cord-Ruwisch et al. 1986).

## Culture medium

Desulfovibrio strain JJ was cultured on medium containing:  $NH_4Cl, 1.0 g; K_2HPO_4, 0.3 g; MgCl_2 \cdot 6 H_2O, 0.2 g;$ L-cysteine · HCl; 0.5 g; CH<sub>3</sub>COONa ·  $3 H_2O$ , 1.0 g; Na<sub>2</sub>SO<sub>4</sub>, 3.0 g; fructose, 4.0 g; yeast extract (Difco Laboratories, Detroit, MI, USA), 1.0 g; resazurin, 0.001 g; mineral solution no. 2 (Balch et al. 1979), 50 ml; trace minerals solution (Balch et al. 1979), 5 ml; destilled water, 1,000 ml. The medium was adjusted to pH 7.0 with 10 M KOH, then boiled under a stream of  $O_2$ -free  $N_2$ -CO<sub>2</sub> (80-20%) and cooled to room temperature. 5 ml of medium were distributed into Hungate tubes using the anaerobic technique of Hungate (1969). After autoclaving (120°C for 20 min), 0.05 ml of 2% Na<sub>2</sub>S · 9 H<sub>2</sub>O and 0.2 ml of 10% NaHCO<sub>3</sub> (sterile, anaerobic solutions) and 0.05 ml of filter-sterilized vitamin solution (Pfennig 1978) were injected into tubes before inoculation. For roll tubes media, 2% agar (Difco) was added.

#### Analytical techniques

Volatile fatty acids, organic acids and alcohols were analysed as previously described (Cord-Ruwisch et al. 1986). Sulfide was measured spectrophotometrically as colloidal CuS (Cord-Ruwisch 1985). Bacterial growth was quantified by measuring the optical density at 580 nm.

### Cell fractionation

The cells (5 g wet weight) were suspended in 10 ml of 50 mM potassium phosphate buffer (pH 7.0) containing 1 mg pancreatic deoxyribonuclease I and disrupted by passing them twice through a French pressure cell (Aminco) at 1,000 atm. The broken cell suspension was centrifugated at 25,000 × g for 20 min to remove cell debris. The resulting cell-free extract was separated into a supernatant and a particulate fraction by centrifugation at 140,000 × g for 2 h. The supernatant was considered as the soluble fraction. The dark gelatinous pellet was resuspended in the same buffer and represented the particulate fraction. Cell-free extracts

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Fig. 1. Phase contrast photomicrograph of *Desulfovibrio fructoso*vorans. Bar is 10 µm

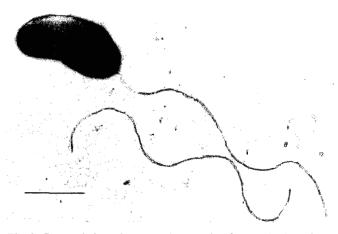


Fig. 2. Transmission electron micrograph of *Desulfovibrio fruc*tosovorans. Note the polar flagellum. Bar is  $2 \mu m$ 

were examined for cytochrome and desulfoviridin using a Cary 219 recording spectrophotometer. Protein was estimated by the method of Lowry et al. (1951) with bovine serum albumin as the standard.

# DNA extraction

Whole-cell DNA was extracted after disruption of the cells and purified according to Marmur (1961) at the German Collection of Microorganisms (DSM), Göttingen, FRG; determination of mol% G+C of the DNA was made by buoyant density centrifugation in a CsCl<sub>2</sub> gradient.

## Results

## Morphology

Greyish round colonies appeared after 1 week of incubation in agar roll tubes at 37°C. The isolate was a vibroid rod (Fig. 1) motile by a single polar flagellum (Fig. 2). It became spirilloid in old cultures. The cells were  $0.5-0.7 \mu m$  in diameter and  $2-4 \mu m$  in length. They occurred singly or in pairs. The cells stained Gram-negative. Spores were never observed.

 Table 1. Utilization of various substrates as electron donors by

 Desulfovibrio fructosovorans in presence and in absence of sulfate

Substrate <sup>a</sup>	$+ SO_4^2$	$- SO_4^2 \sim$
$H_2 + CO_2$	+	
Formate	+	
Ethanol	(+)	_
Glycerol	+	+
Lactate	+	
Pyruvate	+	+
Malate	+	+
Fumarate	+	+
Fructose	+	+

+: Substrates used as electron donors with increase of turbidity (+): Substrates used as electron donors without increase of turbidity <sup>a</sup> Substrates were tested at concentrations of 20 mM. Yeast extract and peptone were tested at concentration of 2 g/l. We also tested the following compounds which did not support sulfide production: acetate, propionate, butyrate, valerate, glutamate, succinate, oxamate, oxalate, choline, ribose, cellobiose, mannitol, inositol, xylose, mannose, inuline, melibiose, maltose, sucrose, galactose, arabinose, rhamnose, glucose, peptone, cysteine, lysine, tyrosine, phenylalanine, asparagine, glycine, tryptophan, arginine, ornithine, alanine. Growth was measured by determining H<sub>2</sub>S or absorbancy after 7 days at 35°C

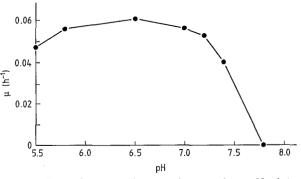


Fig. 3. Effect of pH on the growth rate of *Desulfovibrio fruc*tosovorans cultivated on fructose. Cultures were incubated at 35°C

# Substrate and optimal conditions

Nutritional studies were performed at 35°C. Growth occurred in the absence of vitamins in the described medium as well as in medium for fatty acid-degrading sulfate-reducing bacteria (Widdel and Pfennig 1984). Elemental sulfur, sulfate, thiosulfate and sulfite served as electron acceptors. The isolate did not reduce nitrate. The compounds used as electron donors in sulfate containing medium are summarized in Table 1. No growth was observed on other sugars or amino acids. Growth with molecular hydrogen or formate required acetate as carbon source. In the presence of sulfate, ethanol, glycerol, lactate and pyruvate were incompletely oxidized to acetate and CO2. Methanol was degraded after the cells had been grown on pyruvate but could not be used as the only energy source, even in the presence of acetate. Formate and methanol were oxidized to  $CO_2$ .

Desulfovibrio strain JJ oxidized fructose, fumarate, pyruvate, malate and glycerol in the absence of sulfate. Fructose was fermented to succinate and acetate; small amounts of ethanol (2 mM) was also produced. Fumarate and malate

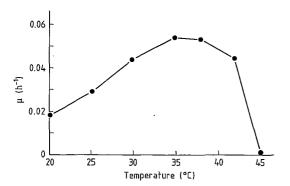


Fig. 4. Effect of temperature on the growth rate of *Desulfovibrio* fructosovorans cultivated on fructose. Cultures were incubated at  $35^{\circ}C$ 

were fermented 'to succinate and acetate. Pyruvate was fermented to mainly acetate. Fermentation products from glycerol were 3-hydroxypropionate and 1,3-propanediol. The isolate grew optimally in culture medium without additionally added NaCl. No growth occurred when 40 g of NaCl per liter was added. The pH optimum for growth was between 6.5 and 7.0 (Fig. 3). The optimum growth temperature was  $35^{\circ}$ C (Fig. 4). In the presence of both sulfate and fumarate, fumarate was the preferred electron acceptor; sulfide was not produced.

## **Pigments**

The soluble extract of *Desulfovibrio* strain JJ exhibited the characteristic absorption bands of cytochrome  $c_3$  with maxima at 418.5, 523 and 552.5 nm when reduced with sodium dithionite. The oxidized extract showed an absorption maximum at 409 nm. The cytochrome was not reduced by sodium ascorbate which indicated that it had a low midpoint redox potential. The spectrum showed in addition a strong absorption band at 628 nm and a weaker one centered at 580 nm characteristic of desulfoviridin (Postgate 1956; Lee and Peck 1971). Fluorescence test for desulfoviridin (Postgate 1959) of cell suspensions was positive.

#### DNA base composition

The mol % G+C content of DNA of strain JJ was 64.13 (mean value of three determinations).

## Discussion

Strain JJ is a curved rod able to reduce sulfate which does not form spores. The isolate oxidizes incompletely pyruvate and lactate to acetate + CO<sub>2</sub>. Neither acetate, propionate nor butyrate are oxidized by strain JJ. Based on these characteristics, the isolate can be affiliated to the genus *Desulfovibrio* (Pfennig et al. 1981). Like in other strains of this genus, the soluble extract of strain JJ exhibits the characteristic absorptions bands of cytochrome c and desulfoviridin (Postgate 1956; Widdel and Pfennig 1984).

The isolate does not belong to the *sapovorans* group since it does not use saturated fatty acids. Morphological and physiological characteristics of strain JJ can be attributed to species *Desulfovibrio desulfuricans* and *D. vulgaris* (Widdel and Pfennig 1984) or to the newly described *D. carbinolicus*  (Nanninga and Gottschal 1986). It differs from *D. desulfuricans* by its ability to oxidize fructose and glycerol and its inability to use choline. In contrast with *D. vulgaris*, strain JJ is able to ferment fumarate and malate. The isolate differs from *D. carbinolicus* in the following ways: it does not use succinate; in the presence of sulfate, glycerol is converted to acetate and  $H_2S$ ; methanol cannot be used as the only energy source, even in the presence of acetate; the strain is motile. A characteristic which distinguishes strain JJ from the methanol oxidizing sulfate reducing bacterium isolated by Braun and Stolp (1985) is that it does not use peptones.

The most important difference between the isolate and all these described species is that strain JJ is the only *Desulfovibrio* strain known that uses a carbohydrate. In the presence of sulfate, fructose is converted to acetate and  $H_2S$ (Cord-Ruwisch et al. 1986). Beside sulfate, also thiosulfate, sulfite, elemental sulfur and fumarate serve as external electron acceptors. Organic substrates are also used in association with hydrogenophilic bacteria acting as electron acceptors (Cord-Ruwisch et al. 1986).

In the absence of sulfate, strain JJ metabolizes not only fructose but also fumarate and malate. Growth on fumarate (Miller and Wakerley 1966) and malate (Miller et al. 1970) was already reported for various *Desulfovibrio* strains. For each fumarate oxidized to acetate via malate and pyruvate, two fumarate molecules are reduced to succinate (Hatchikian and Le Gall 1970a, b; Hatchikian 1972). Succinate is also an end product of fructose fermentation. These results suggest that fumarate is probably an intermediary product and is used as electron acceptor during the fructose degradation. Strain JJ produces small amounts of ethanol during fructose fermentation. Similar results were obtained from fructose using homoacetogenic bacteria (Möller et al. 1984).

Sulfate-reducers are involved in the anaerobic microbial decomposition of organic matter and several results point out the fact that it might exist other nutritional types of sulfate-reducing bacteria in anaerobic environments (Jorgensen and Frenchel 1974; Pfennig et al. 1981). In 1980, Widdel described new morphological types of SRB. Braun and Stolp (1985) and Nanninga and Gottschal (1986) reported on the degradation of methanol by *Desulfovibrio* strains. Utilisation of fructose by strain JJ extends the range of substrates metabolized by this genus. The isolate is proposed as a new species in the genus *Desulfovibrio*: *Desulfovibrio vibrio fructosovorans*.

## Description of Desulfovibrio fructosovorans species nova

*Desulfovibrio fructosovorans* sp. nov.; fruc.to.so.vo.rans. E.m.fructose; L.V.*voro* to devour; M.L.part.adj.*fructoso-vorans* devouring fructose.

Morphology. The cells are curved rods, motile by a single polar flagellum,  $0.5-0.7 \,\mu\text{m}$  wide and  $2.0-4.0 \,\mu\text{m}$  long, occurring singly or in pairs and becoming spirilloid in old cultures. Cells are Gram-negative and no spores were observed.

*Physiology*. Optimum growth occurs at 35°C. The optimum pH for growth is between 6.5 and 7.0.

*Nutrition.* Strictly anaerobic. Elemental sulfur, sulfate, sulfite and thiosulfate serve as electron acceptors and are

reduced to  $H_2S$ . Nitrate is not reduced. Molecular hydrogen, formate, lactate, pyruvate, glycerol, fumarate, malate and fructose serve as electron donors. Growth with  $H_2$  and formate requires acetate as carbon source. Methanol is degraded after the cells have been grown on pyruvate. Fructose, pyruvate, malate, fumarate and glycerol are used in absence of sulfate. Not used: acetate, propionate, butyrate, succinate, oxamate, oxalate, choline. Vitamins are not required for growth. Sodium chloride is not required and is inhibitory above 4% (w/v).

*Pigments*. Presence of desulfoviridin and cytochrome  $c_3$ .

DNA base ratio.  $64.13 \mod \% G + C$ .

*Type strain.* Strain JJ is deposited in the German Collection of Microorganisms, Göttingen, FRG under the collection number DSM 3604.

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