Sulphur-oxidising bacteria and haemoglobin in gills of the bivalve mollusc *Myrtea spinifera*

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ABSTRACT: The gills of Myrtea spinifera carry large numbers of Gram-negative bacteria within the epithelial cells of the subfilamentar region and contain elemental sulphur. Haemoglobin which has a high affinity for oxygen is present in the gill tissue. Homogenates of gill tissue fix carbon dioxide via ribulosebisphosphate carboxylase and contain 3 enzymes concerned in the oxidation of sulphur, adenylylsulphate reductase, sulphate adenylyltransferase and sulphate adenylyltransferase (ADP). Fixation of carbon dioxide by whole gills is enhanced by free sulphide. Whole gills accumulate labelled sulphur when supplied with Na235S, and a major proportion of the labelled sulphur is metabolized to non-volatile compounds or is bound to the tissues. The bacteria are evidently chemoautotrophs and presumably obtain energy for fixation of carbon dioxide by oxidation of reduced sulphur compounds. The haemoglobin is not apparently involved in transport of sulphide but may control oxygen tension close to the bacteria. The sediment in which the bivalves live is low in sulphide compared with some coastal muds and the hydrothermal vents. The interstitial water contains less than $1~\mu M$ of dissolved sulphide and the sediment releases less than 700 $\mu moles~dm^{-3}$ of dissolved sulphide when treated with dilute acid. Habitats with similar low concentrations of dissolved sulphide are widespread in nature. An ability to exploit such low levels of reduced sulphur must be mutually advantageous to host and bacteria, though as yet we have no direct evidence for transfer of organic matter from bacteria to host. To obtain sufficient energy for growth and maintenance of the association, the bacteria must have access to the sediment-bound sulphide.

INTRODUCTION

Sulphur-oxidising bacteria are widespread in sediments and waters where oxic and anoxic conditions meet (Jørgensen 1977, 1982). They obtain energy by oxidation of reduced sulphur compounds and fix carbon dioxide to form organic carbon compounds. A new site for such activity was found in deep-sea hydrothermal vents in 1977. The vent waters contain high concentrations of H₂S and support growth of sulphuroxidising bacteria (Jannasch & Wirsen 1979). The dense communities of animals which surround some of the vents were at first assumed to feed on bacteria suspended in the vent water, but it soon appeared that one of the most abundant creatures was a large gutless worm-like animal (Riftia pachyptila Jones) which cannot take in particulate food. This animal contained vast numbers of endosymbiotic bacteria, probably sulphide oxidizers, which contributed to the nutrition of their host (Cavanaugh et al. 1981). Two large-sized species of bivalve mollusc, abundant around vents, have also

been found to carry similar bacteria in their gills. The bacteria may contribute to the bivalves' nutrition, although these animals do have a normal filter feeding mechanism and a digestive system (Felbeck et al. 1981, Cavanaugh 1983). Animals living in other marine environments with high concentrations of free sulphide, such as sewage outfalls and seagrass beds, also contain symbiotic sulphur bacteria. As a consequence it has been assumed that a fairly high concentration of sulphide is a prerequisite for development of such an association. However, there are other gutless worms (Pogonophora), related to Riftia, containing symbiotic bacteria capable of fixing CO₂ (Southward et al. 1981, Southward 1982). Free sulphide is barely detectable in the sediments inhabited by these pogonophores and other energy sources, including methane, hydrogen and ammonia, have been investigated in the search for their method of nutrition.

A clue to the nutrition of these pogonophores may be obtained from a study of other kinds of animals in the same habitat. In particular, the bivalve superfamily

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Lucinacea was suggested (C. M. Cavanaugh pers. comm.) as a probable source of animals with endosymbiotic bacteria. This superfamily contains a number of species that are typically found in the anoxic layer of sediments and which display unusual gill, palp and alimentary canal structures that have puzzled previous investigators of their functional anatomy (Allen 1958). Recently it has been found that 2 lucinaceans, Lucinoma annulata and Codakia orbicularis, carry endosymbiotic bacteria in their gills, comparable to those also found in the protobranch bivalve Solemya velum (Cavanaugh 1983, Berg & Alatolo 1984).

Three lucinaceans, Lucinoma borealis (L.), Myrtea spinifera (Montagu) and Thyasira flexuosa (Montagu), co-occur with a rich population of the pogonophore Siboglinum fiordicum Webb in a Norwegian fjord (Southward et al. 1979). We have now found that all 3 bivalves contain bacterial symbionts. The bivalves are more convenient to study than small pogonophores. This paper describes the symbiosis in M. spinifera, including the way in which the association appears to be able to exploit the very low concentration of free sulphide in its habitat. Other papers will deal with L. borealis, T. flexuosa and S. fiordicum. It is of great interest to discover such a capability, and to consider the possibility that the very widespread low sulphide environments in the sea (Volkov & Rozanov 1983) may be utilized by similar symbiotic associations between sulphide-oxidizing bacteria and animals.

MATERIAL AND METHODS

We have studied a rich population of Myrtea spinifera (shell length 6 to 12 mm) living in muddy sand at 33 m depth in Ypsesund, a side-branch of Herdlafjorden, near Bergen (Norway), where densities can reach 25 m⁻². Other bivalves present include *Lucinoma* borealis, Thyasira flexuosa, Arctica islandica (L.) and Corbula gibba (Olivi). Specimens were collected with a modified Van Veen grab and kept in muddy sand from the habitat under the cool flowing circulation water at the Institutt for Marinbiologi, University of Bergen. A number of specimens were brought back to Plymouth (England) and kept in a constant-temperature room at 8 °C. Observations were made on the burrowing activity of the bivalves in test tubes filled with sediment. Dissected material was assessed for gill colour and appearance.

Sediment sampling. A gravity corer, fitted with 60 mm internal diameter PVC liners, was used to collect sediment samples. The cores were sealed with bungs as soon as possible after retrieval. Surface samples of sediment were taken through the top flap of the Van Veen grab as soon as it reached the deck. Cores were handled in a glove bag in an atmosphere of either

oxygen-free nitrogen or helium. Interstitial water was collected either by pressure filtration, under nitrogen, through 0.2 µm cellulose acetate filters, or by centrifugation after addition of 1, 1, 2-trichlorotrifluoroethane (Kinniburgh & Miles 1983). Dissolved sulphide was estimated by the methylene blue technique (Cline 1969), acid reagent being added to the samples immediately after collection. Acid-labile sulphide was fixed and trapped as described by Munson (1977) and estimated colourimetrically by the methylene blue technique. Thiosulphate, trithionate and tetrathionate were determined in pore water using the procedure of Kelly et al. (1969). Sub-samples of the cores (5 cm³) were weighed and dried to constant weight at 110 °C to determine water content and then heated to 450 °C to constant weight to estimate organic content. Iron and manganese were measured by atomic absorption spectroscopy after extraction with cold 10 % HCl. Sediment temperature and a measure of the Redox potential were obtained by pushing probes into freshly extruded cores and Van Veen grab samples. A Radiometer oxygen electrode was used in conjunction with an M.B.A. portable oxygen analyser to provide an approximate measure of oxygen content of the upper few cm of sediment.

Enzyme assay. Tissues were dissected out and rinsed with filtered sea water before homogenization in 100 to $300~\mu l$ of ice-cold distilled water in an all glass Potter-Elvehjem homogenizer. Assays were carried out at 12 °C, the highest temperature the animals are likely to experience at Ypsesund, and buffers were adjusted to this temperature. Adenylylsulphate was prepared enzymatically from sodium sulphate labelled with ³⁵S (Amersham) as described by Tsang et al. (1976). Purity of the preparation was checked by paper electrophoresis at pH 5.8 (Baddiley et al. 1957) and corrections were made for sulphate formed by degradation of the adenylylsulphate ... ³⁵S labelled adenylylsulphate was used for assay of adenylylsulphate reductase (EC. 1.8.99.2) following the procedure of Peck et al. (1965). The reaction medium of 500 µl was 160 mM imidazole, 8 mM EDTA, 0.25 mM β-mercaptoethanol, 4mM methylviologen, 5 mM sodium dithionite and 5 μ M adenylylsulphate (35S labelled), pH 7.0. Incubations were carried out in an atmosphere of helium in Warburg flasks. At the end of the experiment 800 µmoles of sulphuric acid were tipped in from one side arm and 8 µmoles of sodium sulphite from the other. The sulphur dioxide was trapped in 100 µmoles of sodium hydroxide in the centre well. Negligible activity was observed in the absence of methyl viologen.

Sulphate adenylyltransferase (EC. 2.7.7.4) and sulphate adenylyltransferase (ADP) (EC. 2.7.7.5) were assayed in a 140 μ l reaction medium of 70 mM imidazole, 7 mM EDTA, 18 μ M adenylylsulphate

labelled with ^{35}S and either 3 mM sodium pyrophosphate or 3 mM sodium triphosphate, pH 7.5. The reaction was stopped by addition of 50 μ l 0.5 M barium chloride, and 50 μ l 20 mM sodium sulphate was added as a carrier, to precipitate released sulphate. The barium sulphate was removed by centrifuging and aliquots of the supernatant were counted for unchanged adenylylsulphate by liquid scintillation spectrometry. Controls containing enzyme but lacking added phosphate or pyrophosphate, when precipitated, showed counts similar to controls containing phosphate or pyrophosphate but lacking enzyme.

Ribulosebisphosphate carboxylase (EC. 4.1.1.39) was assayed in a 70 µl reaction mixture of 43 mM Tris, 20 mM magnesium chloride, 20 mM reduced glutathione, 82 mM sodium bicarbonate, 0.6 mM NaH14CO3 and 5 mM ribulose bisphosphate, pH 8.0. Oxygen was not excluded during the assay. The reaction was stopped with 200 µl glacial acetic acid. After transferring the mixture to glass vials it was evaporated to dryness at 55 °C, reacidified with 200 μl glacial acetic acid and redried before counting. Before use the ¹⁴C labelled sodium bicarbonate supplied by Amersham was dissolved in 1 mM sodium hydroxide and exposed for 24 h to UV light to remove non volatile 14C compounds. After this treatment, controls incubated without ribulose bisphosphate showed negligible fixation of ¹⁴CO₂.

Elemental sulphur. Gills were fixed in 4% formaldehyde in distilled water containing 2% cadmium chloride. Elemental sulphur was extracted with carbon disulphide and reacted with cyanide before measurement as thiocyanate (Fliermans & Brock 1973).

Fixation of sulphur. Fresh whole gills were incubated in filtered sea water containing approximately $18 \mu M l^{-1}$ sodium sulphide (Amersham SJ. 21, Lot' 4/84, specific activity 16.9 $\mu Ci/\mu M$, purity not specified) of which about 1% was 35S. After incubation the gills were rinsed three times with filtered sea water and extracted in alkaline 80 % methanol. In some of the gills elemental sulphur was dissolved out with carbon disulphide and any 35S counted. Other gills were digested in 'NCS Tissue Solubilizer' and the total 'bound' 35S counted. Aliquots of the methanol extract were counted in 'Instagel'. Other aliquots were subjected to electrophoresis in 1 % triethylamine (pH 12), before or after heating with glacial acetic acid to remove volatile sulphur compunds, or were subjected to thin layer chromatography in isopropanol:formic acid: water (6:1:1 v/v). Labelled compounds were detected by scanning the chromatograms and electrophoretograms with a gas flow counter, by autoradiography and (for electrophoretograms) by counting 5 mm strips by liquid scintillation.

Fixation of bicarbonate in presence of sulphide.

Fresh whole gills were incubated in filtered sea water containing known amounts of NaH14CO3, with and without addition of buffered sodium sulphide solution. A parallel control, without gill, was monitored for dissolved sulphide, and additions of sulphide were made to the experimental vessels to maintain the desired level (ca. 20 to 30 μ M l⁻¹). At the end of the experiment the level of sulphide in the experimental vessels was checked. The gills were washed 3 times in filtered sea water, weighed, and immersed for 1 h in sea water acidified to pH 2.5 with hydrochloric acid, to remove bicarbonate. After another rinse the gills were digested in 'Protosol' at 55 °C, and the resulting solution counted in 'Instagel'.

Protein separation. Gill proteins, including haemoglobin, were extracted from 133.6 mg wet weight of gill tissue removed from 12 Myrtea spinifera. The gills were homogenized in ice-cold buffer, pH 7.5, (50 mM Tris, 100 mM sodium chloride, 50 mM magnesium chloride and 10 mM calcium chloride), with an allglass Potter-Elvehjem homogenizer. The homogenate was centrifuged at 12000 g and the pellet was rehomogenized in the same buffer and centrifuged again. The 2 supernatants were pooled and chromatographed at 2 °C on a 1.3 × 51 cm column of Sephadex G-100, in equlibrium with the extraction buffer. Binding of sulphide to the column fractions was investigated by dialysing 500 μ l aliquots against 10 μ M sodium sulphide labelled with 35S, in the pH 7.5 buffer. After 1 h aliquots were removed from the dialysis sacs for liquid scintillation counting. Oxygen binding properties of the column-purified haemoglobin were measured spectrophotometrically (Benesch et al. 1965). Samples were dialysed overnight versus the appropriate buffer before analysis. Sodium dodecylsulphate (SDS) gel electrophoresis was carried out on 14 % polyacrylamide slabs (Laemmli 1970).

Microscopy. Gills were fixed in 2.5% glutaral-dehyde in 0.1 M phosphate buffer, pH 7.3, with 14% sucrose, and also in 2.5% glutaraldehyde in sea water. Small pieces were postfixed in 1% osmic acid in either phosphate buffer or sea water. All were dehydrated in ethanol series and embedded in Spurr resin. Thin sections were stained with uranyl acetate and lead citrate and examined in a Phillips 300 electron microscope. Semi-thin sections were stained with toluidine blue for light microscopy and fresh pieces of gill were examined by phase contrast light microscopy.

RESULTS

General anatomy and behaviour

Anatomy and behaviour of some lucinids were described by Allen (1953), who noted the ciliary cur-

rents of the gill and mantle of *Myrtea spinifera* but did not describe its burrowing or the histology of the gill. Captured specimens burrow readily, under laboratory conditions, by digging with the foot, in the same way as other lucinids, to reach a depth of 10 to 50 mm below the surface of the sediment. Like other lucinids, *M. spinifera* uses the vermiform tip of the foot to construct a narrow, mucus-lined 'inhalent tube' which allows it to draw water from the surface. The water enters the mantle cavity in front of the anterior adductor muscle (Fig. 1). The inlet tube formed in the laboratory can be

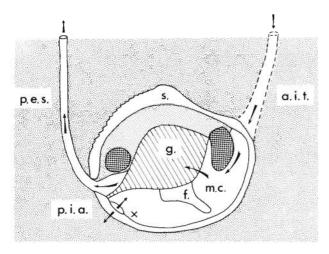


Fig. 1. Myrtea spinifera in sediment (diagrammatic). Arrows show direction of water flow. Dense stipple: adductor muscles; a.i.t.: anterior inhalent tube; f.: foot; g.: gill; m.c.: mantle cavity; p.e.s.: posterior exhalent siphon; p.i.a.: posterior inhalent aperture; s.: shell; x: place where pseudofaeces collect

readily distinguished by its rust-coloured lining of oxidised iron, which contrasts well with the surrounding black, anoxic sediment (Fig. 2). Freshly collected individuals have a similar rust-brown patch at the anterior end of the shell where the inhalent tube adheres. Under laboratory conditions the undisturbed animals form a new inhalent tube, in a different position, every few days. In addition to the inhalent tube of sand grains and mucus, there is a posterior exhalent siphon, with a muscular wall, that can be extended out of the posterior end of the shell and upward to the sediment surface. Below the base of this exhalent siphon, where many bivalves have an inhalent siphon, lucinids have only a small inhalent aperture (Fig. 1). We are not sure yet of the depth at which M. spinifera lives at Ypsesund, since no bivalves were taken in the corer, but if they live at the depths to which they burrow in the laboratory, then the surrounding sediment will have an uncorrected redox reading of $-200\,\mathrm{mV}$, or lower, and oxygen levels will be below those detectable with portable oxygen analysers. Hence any water entering through the posterior inha-



Fig. 2. Myrtea spinifera burrowing in dark muddy sand in a 15 mm diameter test tube; note light colour around inhalent tube, due to oxidation

lent aperture will be pore water from the sediment, containing sulphide and other reduced compounds.

Site of symbionts

When the living shell is opened, the foot and visceral mass can be seen flanked by 2 thick dark red gills (Fig. 3A). The foot is white, but additional red pigment is present in the posterior adductor muscle, the base of the exhalent siphon, and the nerve ganglia. Each of the gills is a single demibranch, not a pair of demibranchs as in most bivalves (Atkins 1937). The gills represent 22.6 \pm 3.6 (S.D.) % of the total body tissue wet weight (11 specimens). A gill of Myrtea spinifera resembles a flat pocket, made up of parallel V-shaped filaments which are connected by rows of interfilamentar bridges to form a pair of grid-like lamellae, joined at the bottom (Fig. 3A, B). The gill lamellae are joined together by plate-like bridges which divide the internal space of the gill into narrow vertical channels which open into a continuous space between gill and visceral mass at the upper end of the gill, extending posteriorly to the base of the exhalent siphon. The main inflow of water to the mantle cavity is through the

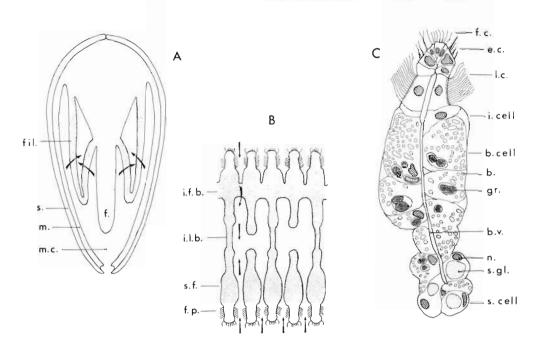


Fig. 3. Myrtea spinifera. (A) Vertical section; arrows: direction of water flow; stipple: position of bacteriocytes. (B) Section through both gill lamellae; arrows: direction of water flow; stipple: position of bacteriocytes. (C) T.S. gill filament. b.: bacterium; b.cell: bacteriocyte; b.v.: blood vessel; e.c.: eulaterofrontal cilia; f.: foot; f.c.: frontal cilia; Fil.: filament; f.p.: filament proper; gr.: brown granule; i.cell: intercalary cell; i.f.b.: interfilamentar bridge; i.l.b.: interlamellar bridge; l.c.: lateral cilia; m.: mantle; m.c.: mantle cavity; n.: nucleus; s.: shell; s.cell: storage cell; s.f.: subfilamentar region; s.gl.: storage globule

anterior inhalent tube (Fig. 1). Water is propelled between the gill filaments by the lateral cilia (Fig. 3C) and flows up the internal channels to the dorsal space (Fig. 3A) and then out via the exhalent siphon (Fig. 1). Particles are filtered off by the cilia on the outer margins of the gill filaments, transported by ciliary currents and sorted so that food particles reach the mouth and rejected particles accumulate as pseudofaeces close to the posterior inhalent aperture (Allen 1958). Any water entering through this aperture will mix in the mantle cavity with the main anterior inflow and pass through the gill. Allen considered that the chief function of the posterior inhalent aperture would be the intermittent disposal of pseudofaeces in the opposite direction. True faeces are extruded from the anus into the exhalent current above and behind the gill and carried out through the exhalent siphon.

The ciliated outer surface of the gill is transparent and the dark red colour shows through from the inner parts of the filaments. Light microscope examination of slices of fresh gill shows that the epidermal cells of the inner parts of the filaments and the interlamellar bridges are inhabited by large numbers of bacteria. Several kinds of inclusion can be seen: (a) colourless, oval to oblong or rod-shaped bacteria; (b) irregular brown and yellow granules; (c) red pigment in separate globules; and (d) a scattering of minute refractile bodies that give the cells a whitish appearance in reflected light and are particularly abundant in the

areas of gill that look whitish in the freshly opened animal, chiefly at the posterior end of the gill. The bacteria can be examined in more detail where damaged cells have released their contents. At least some of the refractile bodies can be seen to be within the bacteria.

From this description it can be inferred that the bacteria-containing cells (bacteriocytes) almost completely line the internal spaces of the gill, and will be exposed to a regular current of water, from which particles have been filtered, whenever the animal is pumping water through the gill (Fig. 3A, B). When the inhalent tube and/or the exhalent siphon is closed, they will be in a stagnant situation, with decreasing oxygen.

Fine structure of gill filaments

The outer ciliated region, or gill filament proper, is similar to that of other bivalves; only the sub-filamentar region is specialized. The epidermis of this region contains 3 types of cell: bacteriocytes, intercalary cells and storage cells. Apparently, mucus cells are absent. The bacteriocytes are the largest and most abundant of the cell-types (Fig. 3C). They are roughly cubical, with a broad base and an outer surface covered with short microvilli. Their cytoplasm is uniformly fine-grained and contains numerous vacuoles enclosing single bacteria (occasionally 2) (Fig. 4). Some larger vacuoles

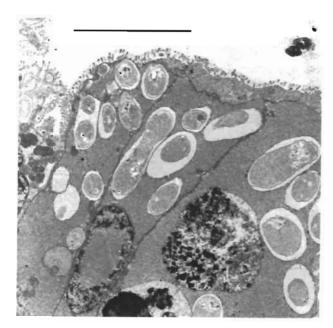


Fig. 4. Myrtea spinifera. Transmission electron micrograph of transverse section of gill filament, showing bacteriocytes partly covered by thin extension of intercalary cell. Note: bacteria in vacuoles; bacteriocyte nucleus, lower left; brown granules, lower centre. Scale bar = $10\,\mu m$

contain varied electron-dense material, which tends to break on cutting and may be rather hard. This corresponds to the brown and yellow granules. Scarce, small, mitochondria lie close to the cell borders. The basal nuclei are unusual in having a large central area clear of chromatin. Possibly this is the location of the haemoglobin concentration seen in living cells (cf. vertebrate erythrocytes, Fawcett 1966). Intercalary cells occur among the bacteriocytes; they are narrower and less electron-dense than the bacteriocytes. The base is narrow but the cell spreads out at the surface of the epidermis to form a thin layer partly covering the neighbouring bacteriocytes (Fig. 4). The short microvilli resemble those on the bacteriocytes and all are enveloped by a thin glycocalyx. The cytoplasm contains more mitochondria, small vacuoles and other inclusions than that of the bacteriocytes. The nucleus lies in the outer part of the cell. Storage cells are dropshaped, with narrow base and rounded outer side (Fig. 3C). These have a smooth surface membrane without microvilli. They are filled with large globular inclusions of electron-dense homogeneous material, bounded by a thin membrane. The cytoplasm contains more obvious endoplasmic reticulum and Golgi membranes than the other 2 cell types. The intracellular globules stain deep blue with toluidine blue in semithin resin sections, while fresh globules are colourless and transparent. They are insoluble in water, ethanol or acetone and do not take up lipid stains. It is probable, therefore, that the storage material is protein.

The central blood space of the subfilamentar region contains haemocytes, including phagocytes carrying granular material similar to the brown granules in the bacteriocytes. This may be waste material which could be disposed of via the digestive system as in other bivalves. There are no haemoglobin-containing cells (erythrocytes) in the blood of *Myrtea spinifera*.

Fine structure of bacteria

The best preservation of internal detail of the bacteria was found in gill material fixed in glutaraldehyde in phosphate buffer with sucrose, post fixed in OsO4 (p.87). There is a wide range of bacterial sizes in any bacteriocyte, the largest found are 2.8 \times 9 μm ; the usual range is from about 2.5×6.5 to 0.4×4 µm. More than 1 species of bacterium may be present, but the internal details are so similar in the different sizes that it seems more probable that they represent development stages of 1 species. The bacterial cell-wall is thin, about 20 nm, and 2 dense layers can be seen, an outer layer and a cytoplasmic membrane. Finer details have not been observed, but the general structure corresponds to that described for typical Gram-negative bacteria. The cytoplasmic membrane is extended to surround small spherical or lens-shaped vesicles 0.1 to 0.5 μm in diameter (Fig. 5), which probably represent the



Fig. 5. Myrtea spinifera. Transmission electron micrograph showing bacterial symbionts, various sizes. Note membrane-bound vesicles in most individuals, smaller dark reticulate bodies in 2 on left. Scale bar = $1 \, \mu m$

small refractile bodies seen with the light microscope. Their contents are preserved during glutaraldehyde fixation but lost during dehydration with ethanol, methanol or acetone, and are quite likely to be elemental sulphur. Other membrane-bound bodies with reticulate, electron-dense contents (0.1 to 0.16 µm diameter) sometimes lie close to the 'sulphur' vesicles, but can occur separately. In some of the larger bacteria there are flat electron-dense crystals, in stacks of 2 or 3 (Fig. 5), possibly without a limiting membrane. The densest of these seem to be homogenous, while the less dense have a layered structure. The largest measured are 1.3 µm long, they may be square, but the exact shape is difficult to determine from sections. The bacterial cytoplasm is filled with irregular particles about 30 to 50 nm in diameter, which may be clumped ribosomes or perhaps glycogen particles. Single, denser ribosomes about 20 nm diameter are visible in a few bacteria. A central core of less dense nucleoplasm extends through most of the length of the smaller bacteria, but is restricted to one end or one side of the larger bacteria (Fig. 4 & 5). This region contains electron-dense DNA strands and also very dense spherical particles, 30 to 100 nm diameter, which are probably polyphosphate (volutin) granules. Thin sections tend to break at these granules. Carboxysomes have not been seen and intracytoplasmic membranes are rare, apart from those surrounding the vesicles and the reticulate granules.

Enzyme activities

The activities of ribulosebisphosphate carboxylase, adenylylsulphate reductase, sulphate adenylyltransferase and sulphate adenylyltransferase (ADP) for tissues of *Myrtea spinifera* and for the gills of *Arctica islandica* and *Corbula gibba* are given in Table 1.

Table 2. Myrtea spinifera. Ribulosebisphosphate carboxylase activity in gills compared with spinach leaf, on a wet weight basis

	V _{max} nmoles g ⁻¹ min ⁻¹	K _m mM
M. spinifera gills	60.1	2.59
Spinach 'King of Denmark'	2501	0.98
Spinach 'Matador'	1158	0.88

Table 2 gives some kinetic values. Only 1 enzyme was found in tissues that were free of symbionts: sulphate adenylyltransferase activity was present, in relatively small amounts when compared with the gill activity of M. spinifera, in the gills of A. islandica and C. gibba, and the foot and mantle of M. spinifera. The other 3 enzymes were confined to the gill of M. spinifera. The addition of 10 mM fluoride had no effect on adenylyltransferase activity and this, together with the absence of magnesium from the medium and the linear time course of the reaction, indicates that inorganic pyrophosphatase activity was not involved. Adenylylsulphate reductase activity was also assayed in gill extracts from 4 other species of bivalves, Venerupis pullastra (Montagu), Nuculana minuta (Müller), Chlamys varia (L.) and a species of Abra. In all these species the measured activity was less than 0.3 n moles g⁻¹ gill min⁻¹ and such low activity could be due to external contamination of the gills by sediment bacteria. The activity of ribulosebisphosphate carboxylase from the gill of M. spinifera was 20- to 40-fold less than in strongly-growing spinach leaves.

Haemoglobin

The haemoglobin content of *Myrtea spinifera* gills was estimated as 0.11 mg per individual, or approxi-

Table 1. Enzyme activities in bivalve tissues, as nmoles of substrate converted to product g^{-1} wet weight min⁻¹ (nmoles g^{-1} protein min⁻¹). Values for individuals

Enzyme	Myrtea spinifera		Arctica islandica		Corbula
	gill	foot	mantle	gill	<i>gibba</i> gill
Ribulosebisphosphate carboxylase (CO ₂ fixed)	28.3, 45.5, 46.5°	nd	nd	0	nd
Adenylylsulphate reductase	15.9, 17.8, 43.2 (150, 158, 172)	1.2, 0.4 (0.9, 2.2)	nd	0	0
Sulphate adenylyltransferase	203, 229, 237, 293 (1625, 1077, 1563)	20, 3 (277, 24)	10 (96)	10 (181)	10 (252)
Sulphate adenylyltransferase (ADP)	89, 112, 151 (709, 527, 996)	0	0	0	0

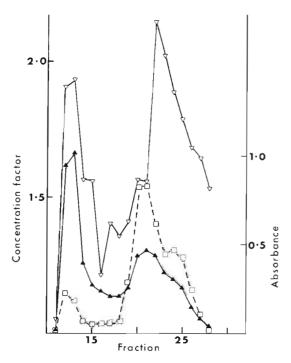


Fig. 6. Myrtea spinifera. Gel filtration of gill extract on Sephadex G-100. Open triangles: sulphide binding (concentration factor) by dialysed fractions; solid triangles: absorbance at 280 nm; open squares: absorbance at 415 nm (haemoglobin), see text

mately 1% of the wet weight of the gills. The pigment separated into 2 fractions on the Sephadex G-100 column (Fig. 6), with an apparent molecular weight (M_r) of 36,300 for the first and 22,000 for the second. When Fraction I was diluted and rechromatographed on the same column, 2 peaks were again resolved with values of 36,000 M_r and 22,000 M_r . SDS gel electrophoresis of the fractions showed that Fraction I consisted of a single 15,450 M_r component (Hb-1), while Fraction II resolved into 3 bands of 24,900, 16,000 and 15,450 M_r . The second fraction would thus appear to contain some fully dissociated Hb-1, together with another haemoglobin, Hb-2. The results from the Sephadex column suggest that the haemoglobin was already dissociating, and if the subunit weight is taken as 15,450 M_r , as

Table 3. *Myrtea spinifera*. Absorption spectra maxima of gill haemoglobin

Protein	λ ^α nm		λ^{β} nm	λ ^{Soret} nm
Oxy Hb Deoxy Hb	581	555	542	415
Carbonyl Hb	571	555	539	432 421

determined for Hb-1 by SDS gel analysis, the associated form would be at least a dimer.

The absorption maxima of the 2 fractions are shown in Table 3 and are similar to those of other haemoglobins. Oxygen binding was determined on very dilute solutions (10 μM) and should therefore be interpreted with caution. At this dilution there was a very high oxygen affinity; at 20 °C the P_{50} was 0.27 torr and the Hill coefficient (n $_{H}$) 1.96, with no detectable Bohr effect between pH 7.52 and 6.75. Fraction I, Fraction II, and a combination of the two in the same proportions found in the gill extract all had similar P_{50} values and cooperativity.

Sulphide binding by gill proteins

Results of dialysis of gill proteins from the Sephadex G-100 column against sodium sulphide are shown in Fig. 6. The concentration of sulphide taken up by the proteins is compared with that present in the 10 μ M dialysis medium and plotted against fraction number alongside total protein (280 nm) and haemoglobin (415 nm).

Sulphide is readily bound by the excluded fraction $(M_r > 100,000)$ and also by a fraction eluting between the 2 haemoglobin peaks, with an estimated M_r of 29,000. There was no obvious correlation between sulphide binding and the haemoglobin fractions.

Sulphur content of gills

Estimates of the elemental sulphur content of the gill of *Myrtea spinifera* are shown in Table 4. There is a wide range from 0.01 to 0.08% of the wet weight, but

Table 4. Myrtea spinifera. Elemental sulphur in gills

Date	Animal condition	Gill colour	Gill weight (mg)	Sulphur (µmoles g ⁻¹)
5. 4. 84 \	Partially starved; kept	Dark red	5.4	5.8
5. 4. 84 ∫	4 wk in laboratory	Dark red	4.9	6.4
22. 6. 84		Deep pink	13	25.0
22. 6. 84	Fixed within 12 h of	Deep pink	15	21.4
22. 6. 84	collection	Dark red	7	16.5
22. 6. 84		Purple red	14	3.3

the higher values were found only in freshly-collected specimens. There is a distinct correlation between colour and sulphur content, the lighter-coloured gills having the most sulphur. We assume that elemental sulphur could be a form of energy storage and indicate 'condition' of the animal/bacteria symbiosis.

Sulphide uptake by gills

Some results for uptake of 35S labelled sulphide by gills of Myrtea spinifera and Arctica islandica are shown in Table 5. The apparent rate of fixation was lower in the longer-term experiments, in which it was likely that the concentration of sulphide would have been reduced by uptake and oxidation. Nevertheless, in *M. spinifera* gills the proportion bound to the tissues increased with time, and a small quantity of it was converted to elemental sulphur. The alkaline-buffer electrophoretograms of the methanol extracts show 2 major peaks of sulphur in M. spinifera but only 1, identified as unchanged sulphide, in A. islandica, while the acid-phase thin layer chromatograms show a larger number of sulphur-labelled compounds in M. spinifera than in A. islandica. Thus gills of both species can take up sulphide, but only M. spinifera, with its

symbionts, is able to metabolize a substantial proportion of the sulphide to other compounds.

Fixation of carbon dioxide

As shown in Table 6, the presence of 30 μ M sulphide increased the total fixation of carbon dioxide by isolated gills of *Myrtea spinifera*, but not by those of *Corbula gibba*. The gills of *M. spinifera* were apparently unchanged after 3 h in 30 μ M sulphide, but those of *C. gibba* shrank considerably and appeared to be damaged by the sulphide.

Sediment chemistry

More details of the chemistry of the Ypsesund sediment will be given in a later paper, but some results that appear relevant to the distribution and biology of *Myrtea spinifera* are given here. For this purpose we shall consider only the top 15 cm of the sediment. The sediment at Ypsesund is a silty fine sand containing shell fragments. The water content varies from 50 to 63 % by volume, and organic matter (by loss of weight on ignition at 450 °C) amounts to 2.8 to 3.2 % of the dry

Gill Time Rate of uptake % uptake % metabolized to still present other compounds wet In 'soluble' form In 'bound' form Total weight as sulphide (% elemental sulphur) $(\mu \text{moles } g^{-1} h^{-1})$ (mg) (min) Myrtea spinifera 0.620.531.15 12 88 (3.5) 8 2 15 8.2 30 0.38 0.36 0.74 26 74 (4.2) 15.7 135 0.15 0.240.39 18 82 (nd) Arctica islandica 45.3 1.5 0.83 1.27 47 53 (nd) 0.44118.2 143 0.33 0.09 34 (nd) 0.41nd: not determined

Table 5. Accumulation and metabolization of 35 sulphide by bivalve gills, at 12 $^{\circ}$ C

Table 6. Fixation of 14 carbon dioxide by bivalve gills exposed to NaH 14 CO $_3$ in seawater, with and without Na $_2$ S (23 to 31 μ M), for 3 h at 12 $^{\circ}$ C

Species	Without su	lphide	With sulphide		
	Fixation rate (nmoles g ⁻¹ min ⁻¹)	Gill wet weight (mg)	Fixation rate (nmoles g ⁻¹ min ⁻¹)	Gill wet weight (mg)	
Myrtea spinifera	0.96	7.2	3.07	6.4	
	0.56	4.7	4.26	3.7	
Corbula gibba	0.32	5.7	0.44	4.8	
	0.75	6.0	0.49	3.7	

weight. Uncorrected redox values become negative a few mm below the sediment surface and range from -200 to -300 mV between 3 cm and 15 cm depth, with a pH between 7 and 8. Sediment temperature (measured after retrieval) was 5.3 to 5.6 °C in February and ca. 8 °C in June 1984. Van Veen grab samples in March showed some oxygen in the near bottom water, but below 5 mm depth in the sediment the oxygen meter read zero.

The level of free sulphide in expressed interstitial water differed according to the method of collection. Two cores from which water was collected by pressure filtration with nitrogen through a $0.2 \mu m$ filter showed no detectable dissolved sulphide in the upper 10 cm. The limit of detection was approximately 1 µM. One core was less than 1 μM , the other gave a value of about 1.5 μM in the 10 to 15 cm depth zone. Higher values for dissolved sulphide were given by interstitial water displaced by trichlorotrifluoroethane, from < 0.5 μM to 4.5 μM for one core, and from 5.7 μM to 11 μM for another. This interstitial water was slightly turbid and, we suspect that the turbidity was partly due to colloidal ferrous sulphide (Volkov & Rozanov 1983) since the most turbid samples usually gave the highest values. Hence the 'dissolved sulphide' values obtained by the Cline method for displaced interstitial water are probably erroneously high. We have obtained free sulphide levels below 1.0 μ M in pore water filtered from similar sediments, and we believe that this value is close to the maximum dissolved sulphide concentration of the interstitial sediment water at Ypsesund. Even this value may be inflated if, as seems likely, finely divided ferrous sulphide passes through the filters used. Additional evidence for the low level of free sulphide can be obtained by olfaction. It has already been reported that the Ypsesund sediment does not smell obviously of sulphide (Southward et al. 1979) and a keen nose is required to detect a very faint smell under field conditions. When such a smell was compared with freshly diluted solutions of sodium sulphide in sea water the limit of olfaction was found to be about 0.1 μM . The smell of 0.5 µM sodium sulphide could be detected quite easily and appeared stronger than that of freshly disturbed sediment.

In contrast to the free sulphide concentration, there was a sharp increase in acid-labile sulphide with depth in the Ypsesund sediment. From zero at the surface this bound sulphide increased to 3.5 $\mu moles$ dm $^{-3}$ of sediment at 1 cm, 315 $\mu moles$ dm $^{-3}$ at 2 cm and 625 $\mu moles$ dm $^{-3}$ at 11 cm. However, the concentration decreased sharply below 25 cm. The concentrations of thiosulphate, trithionate and tetrathionate in the pore water were below, or not significantly different from, the limit of detection of approximately 10 μM .

The concentration of iron, soluble in 10% hydro-

chloric acid, ranged from 80 to 108 m moles dm^{-3} of sediment, and of manganese from 1.2 to 1.4 m moles dm^{-3} . The highest values for both elements were found in the upper 2.5 cm of the sediment.

DISCUSSION

The bacteria in the subfilamentar region of the gill of *Myrtea spinifera* appear to be chemolithotrophic sulphur oxidizers. This conclusion is based on the presence of ribulosebisphosphate carboxylase, adenylyl sulphate reductase and sulphate adenylyltransferase (ADP), together with elemental sulphur, in gill extracts. These three enzymes have not been reported from animal tissues that lack symbiotic bacteria, and were absent from other bivalves living in the Ypsesund sediment, *Corbula gibba* and *Arctica islandica*, which have no symbiotic bacteria in their gills.

Another sulphur-metabolizing enzyme, sulphate adenylyltransferase, was abundant in the gill of M. spinifera, but occurred also in lesser quantities in the foot and mantle tissues which lack symbionts. It was also present in small amounts in the gills of Corbula gibba and Arctica islandica. However, the activity in the gill of M. spinifera was much higher than in the other tissues, suggesting that most of the activity originated in the endosymbiotic bacteria. The presence of elemental sulphur in the gill of M. spinifera also indicates active sulphur metabolism for, although quite small amounts are reported, the analytical method used may underestimate total membrane-bound sulphur: for example, in Thiobacillus concretivorus only 5 to 10% of the sulphur can be extracted by carbon disulphide (Moriarty & Nicholas 1970).

In comparison with known sulphur-oxidizing bacteria, the symbionts from the gills of *Myrtea spinifera* are morphologically closest to the genera *Thiobacterium* and *Thiovulum* (La Rivière & Schmidt 1983). Members of both these genera are oval and spherical and contain numerous sulphur inclusions. *Thiovulum* is a motile slime-producer, while *Thiobacterium* builds up rounded or branching colonies enclosed in slime. A capacity for slime production might be a useful feature in a symbiont required to produce extracellular organic compounds for the nourishment of its host.

Myrtea spinifera gill resembles the gill of Solemya reidi (Felbeck et al. 1981, Felbeck 1983) and the trophosome of Riftia pachyptila (Cavanaugh et al. 1981, Felbeck 1981) in containing bacteria which, from the presence of ribulosebisphosphate carboxylase and adenylylsulphate reductase, are probably chemolithotrophic sulphur oxidizers. There is evidence for this putative symbiosis in other bivalves: Solemya

velum has bacteria in the gill, possesses ribulosebisphosphate carboxylase and shows enhanced fixation of bicarbonate in the presence of free sulphide (Cavanaugh 1983); the gill of Calyptogena pacifica also contains ribulosebisphosphate carboxylase, a high activity of sulphate adenylyltransferase and oxidizes sulphide (Felbeck et al. 1981, Childress & Mickel 1982). Other bivalves that have bacteria in the gill and show ribulosebisphosphate carboxylase activity and high levels of sulphate adenylyltransferase are Calyptogena magnifica and Lucinoma annulata (Felbeck et al. 1981, Cavanaugh 1983) and Codakia orbicularis (Berg & Alatolo 1984). Adenylylsulphate reductase could not be detected in C. orbicularis gill extracts (Berg & Alatolo 1984).

The gutless marine oligochaete *Phallodrilus leukodermatus* has numerous Gram-negative bacteria beneath the cuticle of its epidermis (Giere 1981), shows ribulosebisphosphate carboxylase activity, high activities of sulphate adenylyltransferase and sulphite oxidase, but no detectable adenylylsulphate reductase (Felbeck et al. 1983). This worm lives close to the redox potential discontinuity layer in coral sands and reaches a maximum population density where the free sulphide concentration is 5 to 32 μ M (Giere et al. 1982). However, it ranges from completely oxidized sand to deep in the reducing layer without access to oxygen and may migrate between the layers (Giere et al. 1982, Giere pers. comm.).

The question of access to oxygen brings us to the function of haemoglobin in the gill of Myrtea spinifera. The haemoglobin extracted was heterogenous, with a polymeric fraction, at least dimeric, as well as a monomeric fraction. The polymer undoubtedly dissociated on dilution, but the subunit size for the majority of the material, 15,450 M_r, and for another apparently distinct haemoglobin of 16,100 M_r, are quite similar to those of other haemoglobins. The distribution of haemoglobin in tissues of bivalves appears to be somewhat capricious (Terwilliger & Terwilliger in press), but haemoglobin has already been described from the gill of another lucinid bivalve, Phacoides pectinatus (Lucina jamaicensis), and it is probably present in another member of the family, Codakia orbicularis (Read 1962, 1965, Berg & Alatolo 1984). The gills of the protobranch Solemya velum also contain haemoglobin (Doeller et al. 1983). The haemoglobin in the gill of P. pectinatus closely resembles that from the gill of M. spinifera in that it is heterogeneous with a monomeric fraction of M_r 14,500 and a polymeric fraction which dissociates on dilution. The haemoglobin from P. pectinatus also has high oxygen affinity, $P_{50} = 0.19$ torr at 25 °C, $n_H = 1$, fairly close to the values given here for $M. \ spinifera \ (P_{50} = 0.27 \ torr \ at \ 20 \, ^{\circ}C, \ n_{H} = 1.96).$ Doeller et al. (1983) reported an in situ oxygen affinity

for S. velum haemoglobin of $P_{50}=1$ to 3 torr, and since the haemoglobin of M. spinifera dissociates on dilution, it is possible that its actual in situ affinity for oxygen is less than reported here. Nevertheless, there is no doubt that all the bivalve gill haemoglobins investigated have quite high affinities for oxygen.

It has been suggested that the haemoglobin of the pogonophore Riftia pachyptila may function as a carrier of sulphide (Arp & Childress 1983). However, it seems unlikely that the haemoglobin of Myrtea spinifera has such a function, since we could find no evidence for the binding of sulphide (Fig. 6). Two possible functions remain for the haemoglobin: (1) it could act as a store for oxygen during periods of closure of the inhalent tube and while the animal was burrowing; (2) it could have a function arising from the finding that sulphur-oxidizing bacteria are frequently microaerophilic (Jørgensen 1982) – the haemoglobin might be needed as a buffer to protect the symbionts from excess oxygen which could interfere with their autotrophic processes and lead to spontaneous oxidation of free sulphide. We are thus suggesting that one of the possible functions of haemoglobin in Solemya velum and the 2 lucinids might be comparable to that of leghaemoglobin of the Rhizobium-legume root symbiosis (Wittenberg et al. 1974). The oxygen concentration at the bacteroid surface in legumes is controlled by leghaemoglobin ($P_{50} = 0.55$ torr) to only 10 nM (Bergersen & Goodchild 1973). A further possible function of the haemoglobin, for which we have no data, might be to protect the animal cells from excessive depletion of the oxygen levels when the bacteria are actively oxidizing sulphur compounds.

With the exception of Myrtea spinifera, and apparently, on occasion, Phallodrilus leukodermatus, all the other organisms that possess endosymbiotic bacteria and whose environmental chemistry has been studied, live in habitats with high concentrations of free sulphide, ranging from 160 µM to 25 mM (see Felbeck 1981, 1983 for references). Solemya velum has been reported from eelgrass beds that 'smelled strongly of hydrogen sulphide' (Cavanaugh 1983) and Codakia orbicularis occurs 5 to 10 cm deep in beds of sea-grass Thalassia testudinarium in 'a zone of high sulphide concentrations' (Berg & Alatolo 1984). Our measurements show the upper limit of dissolved sulphide in the sediment at Ypsesund is 1.5 μ M, and the concentration may be as low as $0.5 \mu M$. M. spinifera therefore lives in sediments where the dissolved sulphide concentrations are 2 to 4 orders of magnitude lower than those reported for other habitats containing bivalves with similar chemolithotrophic sulphide-oxidizing symbionts. It seems probable that the low levels of dissolved sulphide in the habitat of M. spinifera are connected with the high concentrations of available iron and manganese. These ions, together with the relatively high pH (for anoxic sediments), may keep the concentrations of sulphydryl ions low (Davison 1980). Nevertheless, the concentration of bound sulphide released by dilute acids, and presumed to be present mostly as ferrous sulphide (Volkov & Rozanov 1983), is considerably lower than reported from other marine sediments (Kaplan et al. 1963, Jørgensen 1977, Novitsky et al. 1980, Troelsen & Jørgensen 1982).

In the absence of data on the rate of sulphate reduction in the Ypsesund sediment we cannot say if the low level of available sulphide is the result of a low rate of production or a high rate of oxidation. The presence of several species of animals with tissues containing symbiotic bacteria will increase the volume of habitat in which sulphide-oxidizing bacteria can exist – they are normally restricted to the oxic/anoxic interface – and could enhance the overall rate of sulphide oxidation in the sediment.

The energy needs of the host animal must be considered. Lucinid bivalves (including Myrtea spinifera, Lucinoma borealis and Codakia orbicularis are filter feeders, but have remarkably inefficient sorting mechanisms, which permit large particles as well as small ones to enter the mouth and reach the stomach. The stomach is simple and the gut short. Allen (1958) concluded that these anatomical modifications were adaptations to a scarce food supply, allowing the use of large particles normally rejected by other bivalves. Uptake of dissolved organic matter (DOM) may play a small part in the nutrition of M. spinifera, but is unlikely to make a substantial contribution unless there is an unexpectedly high concentration of DOM in the inflowing water (Wright 1982). The symbiotic bacteria would be useful if they could supply part or most of the energy needs of the host.

Microbial processes and the chemistry of sulphide oxidation have been reviewed by Jannasch (1983) and Kelly (1982), while Kelly & Kuenen (1984) have calculated the energetics of sulphide-oxidizing symbionts in supplying organic carbon to the host animal. We can extend the calculations of Kelly & Kuenen (1984) to provide estimates of the flux rate required to support the population of Myrtea spinifera. These authors estimated that each mole of sulphur oxidized would yield 0.2 mole of carbon fixed, and they assumed a conversion efficiency of 20%, from bacterium to host. The density of M. spinifera at Ypsesund is 25 m⁻²; the average wet weight of tissue is 65 mg individual⁻¹; we assume that sulphide is drawn from the upper 5 cm of sediment only. Conversion of wet weight to carbon on a 10% basis gives 6.5 mg carbon individual⁻¹, which would correspond to oxidation and conversion of 13.54 mmoles sulphur. If we allow from 2 to 10 yr for growth to adult size then the sulphur utilization rate would be

155 to 773 nmoles h⁻¹. The measured uptake by excised gills in vitro was much lower than this, from 6.0 to 9.4 nmoles h⁻¹, according to incubation time (Table 5), but it is probable that the incubation conditions, including oxygen concentration, were far from optimal. This uptake rate is equivalent to 0.26 μ moles h⁻¹ g⁻¹ body weight and compares with rates of up to 0.5 μ moles h⁻¹ g⁻¹ for *Calyptogena pacifica* (Childress & Mickel 1982) and 1.1 to 1.7 μ moles h⁻¹ g⁻¹ for *Solemya reidi* (Felbeck 1983).

The sediment flux rate required to maintain the estimated needs of 25 Myrtea spinifera m-2 can be calculated as 77 to 386 nmoles sulphide l-1 sediment h^{−1}, not allowing for maintenance metabolism or reproductive effort. If we assume the dissolved sulphide concentration in the environment is $< 1 \mu M$ then adult M. spinifera would have to pump between 150 and 770 ml of interstitial water h^{-1} to obtain sufficient sulphide. These volumes are more than an order of magnitude higher than would be expected from a bivalve of the size of *M. spinifera* (Ola Vahl pers. comm.). It is possible therefore that the free sulphide concentration in the immediate vicinity of the animal may be considerably higher than 1 μ M. The zone of maximum sulphide production in coastal sediments is normally within 5 cm of the surface (Reeburgh 1980, Devol et al. 1984, Troelsen & Jørgensen 1982) and M. spinifera may further raise the production rate in this layer by enriching the neighbouring sediment with a deposit of pseudofaeces (Fig. 1) which might be a nutrient source for sulphate-reducing bacteria. The sediment around the shell is black, apart from the oxidized layer around the inlet tube. Another possibility is suggested by the observation that M. spinifera constructs a new inlet every few days. The oxygen in the inhalent water appears to oxidize the iron sulphides in the sediment around the tube (Fig. 2). The partial oxidation products formed, for example thiosulphate, could diffuse into the inhalent current and contribute to the energy sources for the endosymbiotic bacteria. The presence of elemental sulphur in the gills of freshly collected animals suggests that the bacteria are normally well supplied with reduced sulphur compounds, and this reinforces the argument for access to concentrations higher than the dissolved sulphide levels measured in the interstitial water.

Myrtea spinifera is widely distributed in shallow sea sediments and in the Fjords (Tebble 1966, pers. obs., communications from A. Fosshagen, P. Johannessen, O. Vahl). In contrast it is uncommon in estuaries and has not been found in the shallow Danish inlets where the sediments contain high levels of sulphide (Jensen & Spärck 1934, Jørgensen 1982). We suggest that the symbiosis represented by M. spinifera and its associated bacteria has developed as an adaptation to life in

low sulphide sediments, which are widely distributed in the natural environment (Volkov & Rozanov 1983). This type of symbiosis could be more widespread than hitherto suspected.

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