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Sulphur signature in the hydrothermal-vent mussel *Bathymodiolus azoricus* from the Mid-Atlantic Ridge

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The sulphur compound composition of the hydrothermal vent mussel *Bathymodiolus azoricus* (Bivalvia: Mytilidae) was analysed in order to determine the specific biochemical characteristics of a thiotrophic mode of nutrition. In specimens collected from two contrasting vent-fields on the Mid-Atlantic Ridge (Menez Gwen and Lucky Strike), substantial differences in the sulphur composition were observed between tissues and sampling sites. Total sulphur content was higher for samples from Lucky Strike than for those from Menez Gwen. The amount of elemental sulphur in the gill of *B. azoricus* was in the same range as values previously reported for the vesicomid clam *Calyptogena phaseoliformis* and the lucinid clam *Lucinoma annulata*. Rings of orthorhombic sulphur or compounds such as polythionates or organopolysulphides excreted in the cytoplasm by the symbionts may account for the large amounts of elemental sulphur evidenced for the first time in an hydrothermal vent mussel. A large proportion of the tissue-sulphur was incorporated into free amino compounds such as taurine, hypotaurine, thiotaurine and cysteine, and in a lesser extent to the tripeptide glutathione. In mantles, sulphur seems to be contained mainly in organic compounds such as proteins, mucopolysaccharides and lipids. However, the occurrence of specific compounds such as thiotaurine and elemental sulphur in gills indicate that in this tissue a large proportion of the sulphur might be involved in specific pathways related to transport, storage and detoxification of sulphide. Moreover, our results suggest a greater reliance on thiotrophy of mussels from Lucky Strike as compared to specimens from Menez Gwen.

INTRODUCTION

Endosymbiosis offers a nutritional advantage to metazoan species in deep-sea hydrothermal vents allowing them to exploit geothermal energy sources (sulphide and methane). Vestimentiferan tubeworms and vesicomid clams house only thiotrophic sulphur-oxidizing bacterial endosymbionts, which derive their energy from the large amounts of sulphide present in the hydrothermal fluids (Childress & Fisher, 1992). In contrast, bathymodiolid mussels can host either thiotrophic or methanotrophic symbionts or both simultaneously (for review see Fiala-Médioni & Felbeck, 1990; Fisher, 1990). Dual symbioses have been observed in mussels from hydrothermal vents in the Mid-Atlantic Ridge as well as cold seeps in the Barbados accretionary prism and the Gulf of Mexico (Fisher et al., 1993; Distel et al., 1995; Robinson et al., 1998; Pruski et al., 2000). It is postulated that the co-occurrence of two symbiont types provides the host with a greater nutritional flexibility and the possibility to exploit a wider range of ecological niches (Trask et al., 1999). Site-related differences in the relative abundance of thio- and methanotrophic bacteria have been found in one species of mussel from the Mid-Atlantic Ridge (Trask et al., 1999). This suggests that dual symbioses may respond to environmental variations and adjust the relative contribution of both symbiont types.

Whether the association is strictly dependent on thiotrophy or rely on both sulphur-oxidizing and methanotrophic symbionts, the host has to provide large amounts of sulphide to its endosymbionts, while protecting itself against sulphide toxicity (Powell & Somero, 1986; Somero et al., 1989). As sulphide spontaneously reacts with oxygen, and may therefore lose its energetic interest for the symbionts, the host must avoid sulphide interacting with oxygen before being delivered to the symbionts. Different adaptations to sulphide toxicity have been observed in symbiotic species (Powell & Somero, 1986). The most common adaptations are the reversible binding of sulphide in the blood of vestimentiferan tubeworms and vesicomid clams (Arp et al., 1987; Doeller et al., 1988; Childress et al., 1991, 1993) and the oxidation of sulphide to elemental sulphur in vesicomids (Vetter, 1985) or to thiosulphate in the mitochondria of solemyids (O'Brien & Vetter, 1990). Although sulphide metabolism has been studied less in mussels, it appears that no sulphide-binding protein is present in their haemolymph and there is yet no evidence of elemental sulphur, a potential storage compound for reduced sulphur, in their tissues (Fisher et al., 1987).

We quantified the main sulphur compounds in *Bathymodiolus azoricus* Von Cosel, Comtet & Krylova, 1999, a mussel harbouring a dual symbiosis, in order to determine the specific characteristics of a sulphur-based

nutrition and examine whether any biochemical differences, potentially related to the relative contribution of the sulphur-oxidizing symbionts, could be observed in specimens collected from two contrasting sites. Special attention was also given to the putative physiological significance of the different sulphur compounds found in this study.

MATERIALS AND METHODS

Biological samples

Bathymodiolus azoricus was sampled from two hydrothermal vent-fields (Menez Gwen, 37°51'N 32°31'W, 850 m and Lucky Strike, 37°17'N 32°16'W, 1650 m) on the Mid-Atlantic Ridge during the DIVA 2 cruise (1994). Mussels were recovered with the French submersible Nautille (IFREMER) in a temperature-insulated box. Gill and mantle tissues were rapidly dissected, frozen and stored in liquid nitrogen until lyophilization and analysis in the laboratory.

Elemental analyses

Elemental analyses were carried out by the 'Service Central d'Analyses' of CNRS (Montpellier, France). Total sulphur content was measured by inductivity-coupled-plasma atomic emission spectroscopy (ICP-AES) and total carbon content was determined using an elemental analyser.

Elemental sulphur quantification

Elemental sulphur was extracted in chloroform for 20 min (Boulègue, 1978) and the light absorption was measured at 280 nm ($\epsilon=826$)³. The detection limit of this method is less than 10 $\mu\text{mol g}^{-1}$ dry weight (DW) (Boulègue, 1978). This method had been used successfully to determine elemental sulphur levels in *Calyptogena* sp. from deep trenches in subduction zones (Boulègue et al., 1987). However, it is necessary to emphasize that all the sulphur present at a zero oxidation state is extracted and quantified by this method; the discrimination of S⁰ from polysulphides being therefore impossible. Throughout this paper, all sulphur compounds at an oxidation state of zero are thus referred to as elemental sulphur.

Amino acid extraction and quantification

Extraction of free amino acids was performed as described in Pruski et al. (1997). Primary amines including the tripeptide glutathione were separated and quantified by reverse phase high performance liquid chromatography (HPLC) after derivatization with ortho-phthalaldehyde and 2-mercaptoethanol (see Pruski et al., 1998 for a detailed description of the analytical procedure).

Protein quantification

Frozen tissue samples (~300 mg) were homogenized on ice in 1 ml of extraction buffer (10 mM Tris HCl, 0.5 mM sucrose, 0.15 mM KCl, pH 7.6) supplemented with aprotinin (10 $\mu\text{g ml}^{-1}$). Homogenates were sonicated for 2 minutes and centrifuged (6000 g for 10 min at 4°C). Total protein contents of the supernatants were determined with the Bio-Rad protein assay kit using bovine plasma albumin as standard.

Estimation of sulphur content in individual sulphur compounds

Sulphur incorporated in proteins was estimated after protein quantification using a 1% protein sulphur content (average sulphur content of proteins ranges from 0.2–2.5%, Javillier & Bertrand, 1959). The proportion of sulphur in the other compounds was calculated from their respective concentration and the number of sulphur atoms they contain: every compound contains one sulphur atom per molecule, except thiotaurine (two sulphur atoms) and metallothioneins (21 sulphur atoms). The quantity of metallothioneins per gram of dry tissue was deduced from values published in Rousse et al. (1998) using an approximate molecular weight of 15 KDa (two metallothionein isoforms with apparent size of 10 and 20 KDa are present in bathymodiolid mussels, R. Cosson, personal communication).

RESULTS

Total sulphur contents (S_{TOTAL}) and individual sulphur compound concentrations were highly variable reflecting large differences between individuals, tissues and sampling sites (Table 1).

On average, S_{TOTAL} ranged from 1.7–3.1% of dry tissue with a trend for higher values in samples from Lucky

Table 1. Sulphur content and individual concentrations of a range of sulphur compounds in the tissues of the hydrothermal vent mussel *Bathymodiolus azoricus* sampled from the Menez Gwen and Lucky Strike sites in the Mid-Atlantic Ridge. Values are expressed in $\mu\text{mol g}^{-1}$ dry weight as mean \pm standard deviation ($N=5$). Metallothionein concentrations were deduced from values previously published by Rousse et al. (1998).

	S _{TOTAL}	S ⁰	Taurine	Hypotaurine	Thiotaurine	Cysteine	Methionine	Glutathione	Metallothioneins
Menez Gwen									
Gill	655 \pm 343	170 \pm 30	152.9 \pm 26.0	34.8 \pm 9.1	16.3 \pm 19.2	30.5 \pm 10.6	0.8 \pm 0.3	2.0 \pm 0.4	0.255
Mantle	530 \pm 156	0	82.7 \pm 11.9	38.3 \pm 11.9	2.0 \pm 0.6	29.5 \pm 4.2	0.7 \pm 0.4	1.4 \pm 0.8	0.056
Lucky Strike									
Gill	967 \pm 374	100 \pm 30	96.0 \pm 15.0	28.8 \pm 10.7	62.1 \pm 54.6	12.9 \pm 1.9	1.4 \pm 0.7	2.2 \pm 0.3	0.228
Mantle	717 \pm 405	0	49.1 \pm 13.4	37.2 \pm 5.8	4.1 \pm 2.7	15.2 \pm 4.2	0.7 \pm 0.2	1.8 \pm 0.4	0.099

Table 2. C/S ratios of gill and mantle tissues of *Bathymodiolus azoricus* and six other deep-sea symbiotic bivalves from hydrothermal vents and cold seeps.

	C/S gill	C/S mantle
<i>Bathymodiolus azoricus</i>		
Menez Gwen	18.7 ^a	26.7 ^a
Lucky Strike	12.2 ^a	18.1 ^a
<i>Bathymodiolus thermophilus</i>	17 ^b	34.6 ^b
<i>Calyptogena magnifica</i>	7.3 ^b	36.5 ^b
<i>Calyptogena phaseoliformis</i>	11.2 ^c	67 ^c
<i>Vesicomya</i> sp. (Barbados)	3.3 ^b	39.7 ^b
<i>Bathymodiolus boomerang</i>	29.5 ^b	48.1 ^b
<i>Bathymodiolus</i> n. sp. (Barbados)	34.3 ^b	98.2 ^b

B. thermophilus, *C. magnifica*, *C. phaseoliformis* and *Vesicomya* sp. house sulphur-oxidizing symbionts (Fiala-Médioni & Métivier 1986, Fiala-Médioni & Le Pennec 1988, Fisher et al., 1994, Nelson et al., 1995, Pruski et al., 2000); *B. boomerang* houses sulphur-oxidizing and methanotrophic symbionts (Pruski et al., 2000) and *Bathymodiolus* n. sp. houses methanotrophic symbionts (Pruski et al., 2000). ^a, this study; ^b, J. Boulègue, unpublished results; ^c, J. Boulègue et al., 1987.

Strike than in those from Menez Gwen (Table 1). S_{TOTAL} was typically higher in the symbiont-containing tissues (Table 1).

Values of the carbon/sulphur (C/S) atom ratio were lower in gills than in mantles (Table 2). Gill C/S ratios in *Bathymodiolus azoricus* were similar to those found in the thiotrophic mussel *Bathymodiolus thermophilus* Kenk & Wilson, 1985, higher than those of three vesicomyids from hydrothermal vents and cold seeps and lower than those of two *Bathymodiolus* species from the Barbados accretionary prism (i.e. *Bathymodiolus boomerang* Cosel & Olu, 1998 with both thio- and methanotrophic symbionts and *Bathymodiolus* sp. with only methanotrophic ones, Table 2).

Elemental sulphur was observed only in the gill tissue and was substantially more abundant in samples from Menez Gwen as compared to those from Lucky Strike (26% and 10% of S_{TOTAL} for Menez Gwen and Lucky Strike mussels, respectively, Figure 1).

The free sulphur amino acid pool was dominated by taurine (49–153 $\mu\text{mol g}^{-1}$ DW), which was nearly twice as abundant in gills as in mantles, and to a lesser extent by hypotaurine and cysteine (Table 1). Thiotaaurine was only found in large amounts in gills and was significantly more abundant in the Lucky Strike mussels (62.1 and 16.3 $\mu\text{mol g}^{-1}$ DW in samples from Lucky Strike and Menez Gwen respectively, Table 1). Levels of free methionine and glutathione were low and fairly constant among tissues and sampling sites (Table 1).

Metallothioneins only accounted for a small proportion of the tissue-sulphur (Figure 1).

Total protein contents were in the range of those reported for shallow mussels (43 \pm 3% and 55 \pm 5% of DW, respectively for gill and mantle tissues of *B. azoricus*). No site-related difference in the protein content was observed. We estimated the amount of sulphur incorporated into proteins to range 1/7–1/3 of the total sulphur; mantles having typically higher protein sulphur content than gills (Figure 1).

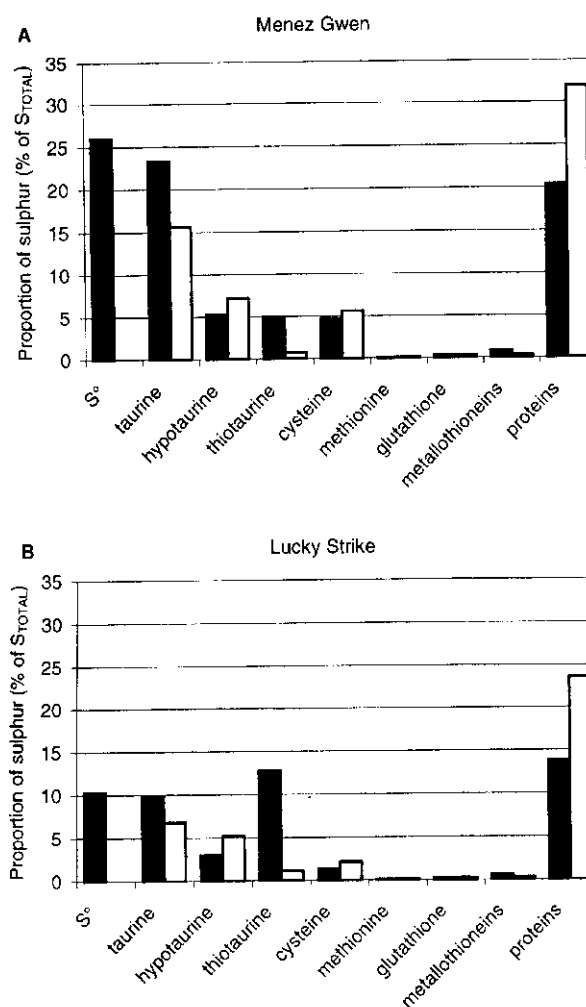


Figure 1. Proportion of sulphur incorporated in a range of sulphur compounds in the tissues of the hydrothermal-vent mussel *Bathymodiolus azoricus* sampled from the Menez Gwen (A) and Lucky Strike (B) sites in the Mid-Atlantic Ridge. Values are expressed as % of S_{TOTAL} (N=5), black bars= gill, white bars= mantle. Every compound contains 1 atom of sulphur per molecule, except thiotaaurine (2 atoms of sulphur) and metallothioneins (21 atoms of sulphur). The proportion of sulphur incorporated to proteins was estimated using a 1% protein sulphur content (average sulphur content of proteins ranges from 0.2–2.5%, Javillier & Bertrand, 1959) and protein contents of 43 and 55% DW, respectively in gill and mantle (see results).

The proportion of unidentified sulphur compounds was higher in mantles than in gills especially in samples from Lucky Strike (respectively in gills and mantles 14% and 38% of S_{TOTAL} in Menez Gwen mussels, and 48 and 61% in Lucky Strike mussels).

DISCUSSION

To date a dual symbiosis has been observed in the two species of *Bathymodiolus* sampled from the Mid-Atlantic Ridge (Fiala-Médioni et al., 1986; Distel et al., 1995; Robinson et al., 1998; Pruski et al., 2000; A. Fiala-Médioni, unpublished data). Both sulphur-oxidizing and methanotrophic bacteria are localized in a single type of host cells (i.e. the bacteriocytes) and contribute to the overall nutrition of their host (Trask et al., 1999).

However, based on the lack of stimulation of carbon fixation by thiosulfate in some species of *Bathymodiolus*, the nutritional importance of the sulphur-oxidizing symbionts in these dual symbioses has been questioned (Cavanaugh et al., 1992). As environmental differences in the chemistry of the surrounding water could potentially modify the relative contribution of the two types of symbionts, and thus their biochemical signature in the tissues, we compared the sulphur compound composition of *B. azoricus* in samples collected from two contrasting sites (Menez Gwen and Lucky Strike).

Total sulphur content in mantles of *B. azoricus* was in the range of those reported for symbiont-free marine organisms, while gills were enriched in sulphur (i.e. sulphur content of marine animals ranges from 0.45 to 2.8%, average values being 0.89%, Kaplan et al., 1963). A high sulphur content in symbiont-containing tissues is a common feature of species relying on thiotrophic endosymbionts for their nutrition (Vetter & Fry, 1998) and can be explained easily by both high environmental sulphur concentrations and the utilization of sulphide by sulphur-oxidizing symbionts. However, a strong enrichment in sulphur was observed in gills of the sulphur-oxidizing symbiont-bearing vesicomyids, *Calyptogena magnifica* Boss & Turner, 1980, *Calyptogena phaseoliformis* Métivier, Okutani & Ohta, 1986 and a non-described species from the Barbados accretionary prism (gill ratios in these three species correspond respectively to 1/5, 1/6 and 1/12 of mantle values). The C/S atom ratio in *B. azoricus* indicates a low sulphur enrichment of the gill tissue compared to the mantle (Table 2). Nevertheless, these values are comparable to those found in the thiotrophic mussel *B. thermophilus* and are therefore consistent with a nutrition based, at least partly, on sulphur-oxidizing endosymbionts (Table 2). Interestingly, gills of vesicomyids always display higher sulphur enrichment than gills of thiotrophic mussels. This may be partly attributable to differences in the chemical composition of their respective environment (e.g. sulphide levels), but it may also be explained by different sulphide uptake mechanisms: vesicomyids actively concentrate sulphide from the environment using a blood component (Childress et al., 1991; 1993), whereas in mussels, which are more passive, sulphide penetrates by simple diffusion (Childress & Fisher, 1992).

Elemental sulphur was found in a wide variety of animals, including several species from hydrothermal vents (for review see Somero et al., 1989; Vetter & Fry, 1998), but was never observed in species of *Bathymodiolus* (e.g. *B. thermophilus* did not contain elemental sulphur, Fisher et al., 1987; Vetter & Fry, 1998). Nevertheless, our results indicate that a large proportion of the gill sulphur is at a zero oxidation state in *B. azoricus* (Table 1), which strongly suggests the occurrence of elemental sulphur (S^0). Those high levels of S^0 are in the range of those reported for thiotrophic species such as the vesicomyid *Calyptogena phaseoliformis* (Boulègue et al., 1987) and the lucinid *Lucinoma annulata* Reeve, 1850 (Vetter, 1985). Microanalyses of *B. thermophilus* gill have only demonstrated the accumulation of sulphur in lysosomes and to a lesser extent in the bacteria (Chassard-Bouchaud et al., 1988). The high amounts of S^0 found in gills of *B. azoricus* may thus represent polar chains of polysulphides that have precipitated in the cytoplasm of the bacteriocytes in the

nanometric orthorhombic form S_8 as suggested for *Riftia pachyptila* Jones, 1981 (Truchet et al., 1998). Other compounds such as polythionates $Sn(SO_3)_2$ - or organopolysulphides $R-Sn-R$, which are commonly produced by sulphur-oxidizing *Thiobacilli* and phototrophic sulphur bacteria (Hazeu et al., 1988; Durand et al., 1994; Prange et al., 1999), could have been extracted and quantified by the method we used. Hence, these compounds could putatively account for some of the elemental sulphur found in *B. azoricus*. However, these compounds have thus far never been observed in thiotrophic symbioses. The amount of S^0 in field specimens represent a balance between exposure to H_2S and metabolic consumption of the resulting S^0 (Vetter & Fry, 1985). As geochemical data indicate that the sulphide level at Menez Gwen is lower than at Lucky Strike (Charlou et al., 1996), the higher elemental sulphur content in the Menez Gwen mussels suggests a lower S^0 consumption, which could reflect a low sulphide-oxidizing activity.

When S^0 content is subtracted from S_{TOTAL} , the amount of sulphur in symbiont-containing tissues are not significantly different from values in mantles, where sulphur is likely to be found in compounds playing no specific function in sulphide metabolism (e.g. free and protein amino acids, sulphated mucopolysaccharides and sulpholipids). Our results indicate that a large proportion of the sulphur was indeed contained in free sulphur amino acids under a reduced form (cysteine, methionine and the sulphhydryl group of thiotaurine) or an oxidized form (taurine, hypotaurine and the inner sulphur of thiotaurine). Taurine was, by far, the dominant sulphur-containing amino acid in all tissues. A high concentration of this compound is a common feature in non-symbiotic bivalves where taurine is one of the main effectors for osmoregulation (Awapara, 1962; Campbell & Bishop, 1970; Bishop et al., 1983). Specific functions related to the symbiotic mode of life have been also attributed to taurine, which could be a sink for sulphur and a nitrogen storage compound as well (Conway & MacDowell Capuzzo, 1992; Lee et al., 1997). The co-occurrence of the two unusual amino acids, hypotaurine and thiotaurine, is a specific feature in deep-sea thiotrophic symbioses (for review see Pruski et al., 2000), where these compounds may be involved in the transfer of reduced sulphur from the host to the symbionts. Moreover, as thiotaurine contains one atom of sulphur at a reduced state ($-II$), it may represent a non-toxic storage compound for reduced sulphur until its oxidation by the sulphur-oxidizing symbionts. It was in fact previously observed in two species of *Bathymodiolus* (*Bathymodiolus brevior* Cosel, Métivier & Hashimoto, 1994 and *Bathymodiolus elongatus* Cosel, Métivier & Hashimoto, 1994) from the Lau Basin that the level of this thio-amine decreased in specimens sampled from declining sites (Pranal et al., 1995). The lower thiotaurine concentration in the gills of the Menez Gwen mussels, together with their higher elemental sulphur content suggest therefore a lower reliance on thiotrophic symbionts in the Menez Gwen mussels. This is consistent with evidence from bacterial counts in gills indicating a higher relative abundance of methanotrophic symbionts in specimens from Menez Gwen as compared to those from Lucky Strike (A. Fiala-Médioni, unpublished data) and geochemical data indicating higher methane content in Menez Gwen fluids (Charlou et al.,

2000). This study also reported higher concentrations of sulphide at Lucky Strike, which can explain the higher S_{TOTAL} values in samples from this site.

Protein sulphur is usually the dominant form of sulphur in animals. It indeed represents an important proportion of the total sulphur content in both tissues (Figure 1). In the mantle, which is involved in the storage of proteins and glycogen (Gabbott, 1983), protein sulphur accounted, as expected, for a larger proportion of the total sulphur content (1/4–1/3 of S_{TOTAL}) than in the gill (1/7–1/5 of S_{TOTAL} , Figure 1). Other organic compounds such as sulphated mucopolysaccharides are typically secreted in the mantle of bivalves (Garcia-Gasca et al., 1994) and could account for a non-negligible proportion of the sulphur in this tissue. Some specific proteins, such as metallothioneins, are known to have a high sulphur content (21 S atoms per molecule, Stillman, 1995) and have been proposed to play a role in thiosulphate transport in *Bathymodiolus* spp. (Rousse, 1999). However, metallothioneins only accounted for a small proportion of the tissue-sulphur in *B. azoricus* and are thus unlikely to be of prime importance for sulphur metabolism (Figure 1).

Considering the specific characteristics of organisms from high sulphide environments, which possess a sulphide-oxidizing peripheral barrier (Powell & Somero, 1986), inorganic oxidized sulphur products ($S_2O_3^{2-}$, SO_4^{2-} , Sn^{2-}) are certainly present in *B. azoricus* tissues. These inorganic sulphur compounds are probably more abundant in gills than in mantles. This was demonstrated for *B. thermophilus* where the lower thiosulphate levels found in the mantle may be explained by the lower activity of the sulphide-oxidizing barrier in this tissue (Fisher et al., 1988).

In summary, most of the sulphur contained in the tissues of *B. azoricus* is incorporated in common organic compounds. However, the sulphur enrichment of the gill of this mussel and the occurrence of specific sulphur compounds, such as elemental sulphur and thiotaurine, are consistent with a nutrition partly based on sulphur-oxidizing symbionts. Our results furthermore suggest that mussels from Lucky Strike are more dependent on thiotrophy as compared to specimens from Menez Gwen.

As *Bathymodiolus*-like species do not possess specific proteins allowing them to concentrate and store sulphur on a non-toxic form, sulphur storage compounds such as elemental sulphur and thiotaurine may be important to support the activity of the bacterial sulphur-oxidizing symbionts during short time periods where sulphide availability in the environment is reduced.

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