

Persephonella atlantica sp. nov.: How to adapt to physico-chemical gradients in high temperature hydrothermal habitats

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| 1 | Persephonella atlantica sp. nov.: How to adapt to physico-chemical |
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| 2 | gradients in high temperature hydrothermal habitats |
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19 Abstract

A novel thermophilic, microaerophilic and anaerobic, hydrogen- sulphur- and thiosulphate-20 oxidising bacterium, designated MO1340^T, was isolated from a deep-sea hydrothermal 21 chimney collected from the Lucky Strike hydrothermal vent field on the Mid-Atlantic Ridge. 22 Cells were short, motile rods of 1.4 - 2.2 µm length and 0.5 - 0.8 µm width. Optimal growth 23 24 was observed for a NaCl concentration of 2.5 % (w/v) at pH 6.5. As for other members of the genus *Persephonella*, strain MO1340^T was strictly chemolithoautotrophic and could oxidise 25 26 hydrogen, elemental sulphur or thiosulphate using oxygen as electron acceptor. Anaerobic 27 nitrate reduction using hydrogen could also be performed. Each catabolic reaction had a 28 different optimal growth temperature (65 to 75 °C) and an optimal dissolved oxygen 29 concentration (11.4 to 119.7 µM at 70 °C for aerobic reactions) that varied according to the 30 electron donors utilised. These experimental results are consistent with the distribution of 31 these catabolic substrates along the temperature gradient observed in active hydrothermal systems. They strongly suggest that this adaptive strategy could confer a selective advantage 32 for strain MO1340^T in the dynamic part of the ecosystem where hot, reduced hydrothermal 33 34 fluid mixes with cold, oxygenated seawater. Phylogenetic analysis indicated that strain MO1340^T was a member of the genus *Persephonella* within the order *Hydrogenothermales* as 35 36 it shared a 16S rRNA gene sequence similarity <95.5 % and ANI respectively 75.66 % with 37 closest described *Persephonella* (*P. hydrogeniphila* 29W^T). On the basis of the physiological 38 and genomic properties of the new isolate, the name Persephonella atlantica sp. nov. is proposed. The type strain is $MO1340^{T}$ (=UBOCC-M-3359^T =JCM 34026^T). 39 40

Keywords 41

42 Aquificota, Persephonella, hydrothermal, chemolithoautotroph, temperature, thermophile

43 Introduction

44 Hydrothermal chimneys are mineral structures resulting from the precipitation of 45 various chemical species during the mixing between the hot, reduced hydrothermal fluid and 46 the cold, oxygenated seawater [31, 34]. These structures and the surrounding seawater are 47 characterized by steep gradients: (1) chemical gradients due to the mixing of reduced compounds (e.g. H₂, CH₄, H₂S, $S_2O_3^{2-}$, Fe²⁺) from the hydrothermal fluid, and oxidised 48 compounds from the surrounding seawater (e.g. O_2 , NO_3^- , SO_4^{2-}); and (2) thermal gradients 49 50 decreasing from the inner parts of chimneys heated by hydrothermal fluid circulation (> 300 51 $^{\circ}$ C), to the external parts in contact with seawater (2-4 $^{\circ}$ C) [45, 51]. 52 These gradients cause a chemical disequilibrium that supports chemoautotrophic 53 microorganisms that oxidise reduced compounds from hydrothermal fluid using oxidised 54 species from seawater. Hence, adaptation of these microorganisms to the thermal and 55 chemical gradients shapes their spatial distribution across hydrothermal habitats [10, 11, 51, 56 74]. For example, dissolved oxygen concentrations are inversely correlated with temperature 57 and usually separate aerobic mesophiles from anaerobic hyperthermophiles [1, 41, 45, 46]. At 58 temperatures exceeding 80 °C, a majority of microorganisms in these environments seem to

have specifically adapted to conditions that are present only at small spatial scales in the hydrothermal chimney (e.g. *Methanococcales, Thermococcales*) [11, 45]. However, at lower temperatures, the diversity of reactants (e.g. oxygen, nitrate, sulphides) seem to favour metabolic versatility as it was described for *Epsilonbacteraeota*, one of the dominant lineages in these habitats [5].

Members of phylum *Aquificota* are another illustration of chemoautrophic
microorganisms metabolically versatile that are ubiquitous in deep-sea hydrothermal vents
[18, 25, 47, 50, 58]. The phylum *Aquificota* (previously known as *Aquificae*) was first
described in 2001 and contains now two classes according to the Genome Taxonomy

68 Database [55], *Desulfurobacteriia* and *Aquificae*. *Aquificae* are divided in two orders: 69 Aquificales and Hydrogenothermales [14, 28, 39, 55]. All members of the phylum Aquificota 70 are thermophilic or hyperthermophilic Gram-negative rods Bacteria, isolated from terrestrial 71 hot ecosystems, coastal, or deep sea hydrothermal vents [29]. They share an obligate or 72 facultative chemolithoautotrophic metabolism by fixing carbon dioxide using reductive 73 tricarboxylic acid cycle (rTCA) [32], with hydrogen or reduced sulphur compounds as 74 electron donors and nitrate as electron acceptors, also oxygen is used as electron acceptor by 75 members of the class Aquificae. Aquificota mostly diverge from other phyla by displaying 76 distinct16S rRNA gene sequences and genomic conserved signature indels [29]. Members of 77 genus *Persephonella*, and other genera in the family *Hydrogenothermaceae* 78 (Hydrogenothermus, Sulfurihydrogenibium and Venenivibrio) share the common traits 79 described above, and show an extensive genomic diversity exhibiting a biogeographic 80 distribution pattern [47]. Although endemic and widely distributed among deep sea 81 hydrothermal vents the genus Persephonella is, to date, only composed of three described species isolated from hydrothermal vents in the Pacific ocean: P. marina EX-H1^T, P. 82 guavmasensis EX-H2^T and *P*. hydrogeniphila 29W^T [25, 50]. 83 84 In this study, we report the isolation of a novel thermophilic, microaerophilic to 85 anaerobic, chemolithoautotrophic *Persephonella* species obtained from an Atlantic deep-sea 86 hydrothermal vent. Characterization of this novel strain demonstrates metabolic adaptation to 87 a wide range of physical and chemical conditions. 88 89 Material and methods 90 Culture, enrichment and isolation 91 The bottom part of the Aisics chimney (sample Chem08; 1690 m depth) [61] was

92 sampled using the ROV Victor 6000 during the MOMAR2008 oceanographic cruise [16] at

93 the Lucky Strike hydrothermal vent field on the Mid-Atlantic Ridge (37° 17' 20.26" N, 32° 94 16' 32.05" W). Chimney sample was brought to the sea surface in a dedicated decontaminated 95 insulated sample box. On board, chimney fragments were crushed under anoxic conditions using an anaerobic chamber and stored at 4 °C in sterile serum vials filled with artificial 96 97 sterile seawater (Sea salts, Sigma Aldrich) under a N₂ atmosphere. Ten years later, 0.5 mL of 98 hydrothermal chimney slurry was used to inoculate the medium described below. X-ray 99 diffraction analysis showed that chimney sample was mainly composed of sulphides including 100 chalcopyrite, pyrite and marcasite (Ewan Pelleter, personal communication). 101 Enrichment cultures were performed in 50 mL glass bottles sealed with butyl septa, 102 and containing 20 mL of the medium described by Parkes *et al.* omitting Na₂S and FeCl₂ 103 solutions [54]. This medium contained per litter: NaCl (24.3 g), MgCl₂.6H₂O (10 g), 104 CaCl₂.2H₂O (1.5 g), KCl (0.66 g). Trace elements were provided by SL-10 solution (1 mL), 105 Selenite-Wolfram solution (0.2 mL), and 1 mL of KBr (0.84 M), H₃BO₃ (0.4 M), SrCl₂ (0.15 106 M), NH₄Cl (0.4 M), KH₂PO₄ (0.04 M) and NaF (0.07 M) solutions. The medium was 107 sterilized by autoclaving for 30 minutes at 100 °C and was buffered by adding NaHCO₃ (2.52 108 g). 109 Thiosulphate was added at a final concentration of 10 mM as electron donor. Gas phase was 110 composed of N₂:CO₂ (80:20, 150 kPa) and 1 mL of sterile air was added with a syringe in 111 order to create microaerophilic conditions. After 12 days incubation at 30 °C without 112 agitation, subcultures from positive enrichment were consecutively incubated 7 days at 50 °C and 7 days at 60 °C. Strain MO1340^T was isolated by three consecutive tenfold dilutions in 113 Hungate anaerobic tubes incubated at 60 °C in the same medium as for the enrichment step. 114 115 Five successive transfers were also performed on medium solidified with 1 % gellan gum 116 (Sigma Aldrich) incubated at 70 °C in serum vials with hydrogen as energy source (gas phase

117 H₂:CO₂ 80:20, 200 kPa) and nitrate as electron acceptor (8 mM). Purity of the strain was

118 regularly checked by microscope observation and 16S rRNA gene sequencing.

119 Morphological characterization

120 Cells morphology and motility of strain MO1340^T were observed under an Olympus
121 BX60 phase contrast microscope. Transmission electron microscopy observations were
122 performed without staining on exponential phase cultures grown on thiosulphate and oxygen
123 at 70 °C (JEM-1400, JEOL; *Plateforme d'Imagerie et de Mesures en Microscopie*, University
124 of Brest).

125

Physiology and growth requirements

126 Unless otherwise stated, all physiological characterizations were done in anaerobic 127 Bellco tubes (Bellco Glass Inc., Vineland N.J.) in triplicate filled with 15 mL of the medium 128 described above at pH 6.8, incubated without agitation at 70 °C. Uninoculated media blanks 129 were used as negative controls for each culture condition. Growth was determined from cell 130 counting by microscope observation using Thoma cell counting chamber (depth: 0.02 mm). 131 Flow cytometry counting was also performed on cells fixed with 2.5 % (w/v) glutaraldehyde and stored at 4 °C. Cell samples were diluted 10 to 100 times in a saline solution with 2.5 %132 133 (w/v) NaCl and stained with SYBRGreen I (Sigma Aldrich). Cells were counted using a 134 Cyflox Space (Sysmex, Partec).

Dissolved oxygen requirement was determined by increasing oxygen concentration in the gas phase from 0 to 21 % at 150 kPa (0 to 179.5 μ M of dissolved oxygen) in 3 different media containing respectively (1) thiosulphate (10 mM); (2) colloidal elemental sulphur 1 % (w/v); or (3) hydrogen (80 % in the gas phase, 200 kPa; 820.7 μ M of dissolved hydrogen) as electron donors. The influence of temperature on growth was determined for temperatures ranging from 40 to 85 °C in 4 different culture conditions: in the presence of (1) thiosulphate (10 mM) under N₂:CO₂:O₂ gas phase (77.5:20:2.5, 150 kPa; 21.4 μ M of dissolved oxygen);

| 142 | (2) elemental sulphur (1 %, w/v), under N ₂ :CO ₂ :O ₂ gas phase (66:20:14, 150 kPa; 119.7 μ M |
|-----|--|
| 143 | of dissolved oxygen); (3) hydrogen in anoxic conditions with nitrate (8 mM) as electron |
| 144 | acceptor under H ₂ :CO ₂ gas phase (80:20, 200 kPa; 820.7 μ M of dissolved hydrogen); and (4) |
| 145 | hydrogen in microaerophilic conditions under H ₂ :CO ₂ :O ₂ gas phase (79:20:1, 200 kPa; 820.7 |
| 146 | μ M of dissolved hydrogen; 11.4 μ M of dissolved oxygen). NaCl concentration and pH |
| 147 | optimum and range were determined on thiosulphate (10 mM) containing medium, under |
| 148 | N ₂ :CO ₂ :O ₂ (77.5:20:2.5, 150 kPa) gas phase. NaCl concentrations tested ranged from 0 to 5.5 |
| 149 | % (w/v). pH was adjusted from 4.5 to 6 by adding increasing volumes of sterile solutions of |
| 150 | 1% (w/v) NaHCO ₃ to medium without buffer. For pH above 6, 2 % (w/v) Na ₂ CO ₃ was added |
| 151 | to medium buffered with 2 mM NaHCO ₃ (instead of 30 mM in standard medium). |
| 152 | Electron donor and acceptor requirement |
| 153 | Electron donors were tested by supplementing mineral medium (artificial seawater |
| 154 | described above, without Na ₂ S nor FeCl ₂) with: hydrogen (80 % in the gas phase, 200 kPa), |
| 155 | formate (10 mM), succinate (10 mM), ferrous iron (10 mM), acetate (1 mM), ammonium |
| 156 | chloride (15 mM) or elemental sulphur (1 $\%$ w/v). Oxygen (1.5 or 7 $\%$ in the gas phase, 150 |
| 157 | kPa) or nitrate (10 mM) were added as electron acceptors in all conditions. The following |
| 158 | electron acceptors were tested in presence of H ₂ :CO ₂ (80:20, 200 kPa): sulphate (10 mM), |
| 159 | sulphite (10 mM), nitrite (1 or 14 mM), ferric iron (10 mM), elemental sulphur (1 % w/v), |
| 160 | acetate (1 mM) or thiosulphate (10 mM). Organic carbon sources were tested in presence of |
| 161 | N ₂ :CO ₂ :O ₂ (73:20:7, 150 kPa) at a final concentration of 2 g/L (yeast extract or peptone) or 5 |
| 162 | g/L (glucose, sucrose, fructose, lactose or sorbitol). Mixotrophic growth was tested using |
| 163 | yeast extract (0.5 g/L), peptone (0.5 g/L) and glucose (1 g/L) as carbon sources and sodium |
| 164 | bicarbonate was replaced by PIPES disodium salt (6 g/L). Hydrogen (200 kPa) and nitrate (10 |
| 165 | mM), or thiosulphate (10 mM) or sulphur (1 $\%$ w/v) in the presence of oxygen (7 $\%$ in the gas |
| 166 | phase, 150 kPa) were provided as energy sources. |

167 Consecutive positive cultures were transferred three times on the same medium to 168 ensure that growth was not due to substrate carry-over from the inoculum. Controls 169 containing no electron acceptor or donor were performed. Utilisation of various electron donors and acceptors were also performed on the closest relative to strain MO1340^T, 170 Persephonella hydrogeniphila 29W^T (DSM 15103) provided by the German Collection of 171 172 Microorganisms and Cell Cultures. Growth in the absence of Balch vitamins solution was 173 tested (as recommended in [71]) using the standard medium in the presence of thiosulphate 174 and oxygen. In order to identify the metabolic reactions performed by strain MO1340, end-175 point cultures were carried out and both substrates consumption and metabolic by-products were quantified. Four couples of substrates were tested $(S_2O_3^{2-}/O_2; S^0/O_2; H_2/O_2; H_2/NO_3^{-})$ in 176 177 the same conditions as described above. At least three negative controls and 3 to 10 culture 178 replicates were performed and incubated at 70 C.

179 *Analytical techniques*

180 Hydrogen, nitrogen and carbon dioxide concentration in the headspace phase were 181 measured using a modified INFICON/Micro GC FUSION Gas Analyser (INFICON, Basel, 182 Switzerland) fitted with a pressure gauge (CTE8005AY0, Sensortechnics GmbH) and two 183 conductivity detectors. Separation was performed using two columns: molecular sieve 10m 184 column and argon as a carrier gas; and a RT-Q 12m using helium as a carrier gas. Gas 185 concentrations were calculated using the method of Mah et al. [42]. Concentrations of 186 dissolved oxygen and hydrogen in the experimental conditions were calculated for seawater at 187 70 °C as described respectively by Weiss et al. [81] and Crozier et al. [9]. 188 Thiosulphate, nitrate and sulphate concentrations were quantified by anion

chromatography using a Dionex ICS-2000 Reagent-Free Ion Chromatography System
(Thermo Fisher Scientific) as described by Webster *et al.* [80]. A potential hydrogen peroxide
production was assayed by using the MyQubit AmplexTM Red Peroxyde Assay (Invitrogen)

according to the manufacturer's recommendation. Stoichiometry of the reactions were
calculated from experimental data and then compared to metabolic reactions described in
literature [2, 78].

195 Thermodynamic calculations and Arrhenius parameters

196 To quantify the energy gained from each reactional pathway considered, we calculated the 197 overall Gibbs free energy of the reaction derived from the following expression (Eq. 1) 198 $\Delta_r G = \Delta_r G_0 + RT ln Q_r$ Eq.1

199 Where $\Delta_r G_o$ represents the standard state Gibbs free energy of reaction, R is the gas constant, 200 T is temperature in Kelvins and Q_r stands for the activity product for the natural conditions. 201 The activity product Q_r is given by Eq.2

202
$$Q_r = \prod a_i^{\gamma_{i,r}}$$
 Eq. 2

203 Where a_i stands for the activity of the *i*th constituent of the reaction r, and $\gamma_{i,r}$ represents the 204 stoichiometric reaction coefficient (negative for reactants, positive for products). Activities 205 are derived from concentrations through the extended Debye-Huckel equation for activity 206 coefficients. Thermodynamic and hydrodynamic modelling were performed on R [77] using 207 the CHNOSZ database [12] and ReacTran [69], gsw [37] and marelac [70] packages. The 208 metabolic pathways and reactions considered in this work are listed in Table 2. 209 Concentrations of the chemical species at various temperatures were determined following 210 two approaches. 1) When available, we used *in situ* data collected along the mixing gradient between end member hydrothermal fluid at Aisics chimney and surrounding seawater using 211 212 the PEPITO water sampler [8], coupled to an oxygen Aanderaa optode and the CHEMINI in 213 situ chemical analyser for total sulphide concentrations [79]. 2) When not available, a simple 214 hydrodynamic model was used to predict dilution processes along the mixing gradient and 215 calculate concentrations assuming simple mixing (i.e. no reaction) between the hydrothermal 216 fluid and the surrounding seawater. Values of the end-member fluid are based on the literature [6, 61] or inferred from other hydrothermal sites when no data are available on Aisics ([19]
for elemental sulphur). *In situ* data were collected during the MOMARSAT 2015 cruise on

219 board R/V Pourquoi Pas? [65]. Fluid samples were analysed back on shore for nitrate,

sulphate and thiosulphate concentrations using anion chromatography as described by

221 Webster *et al.* [80].

222 Concentrations of the end-member fluid might have changed since the strain was initially

223 collected in 2008. However reported concentrations of Aisics high temperature fluids show

limited variability [6, 61] and we therefore expect only minor alteration of the distribution of

225 chemical species along the mixing gradient over the years.

226 The effect of temperature on specific growth rate was studied using Arrhenius plots

and Q₁₀ values that were calculated as described by Roussel *et al.* 2015 [62]. Statistical

analysis and multiple peek fitting were performed using Origin 2016 (OriginLab Corporation,

229 Northampton, USA) with the multiple peak fit tool using the Gauss peak function.

230 *Genome sequencing and analysis*

231 Genomic DNA of strain MO1340^T was isolated by PCI extraction [82] from cells

grown on thiosulphate and oxygen at 70 °C. 16S rRNA gene was amplified by PCR using

bacterial primers E8F and U1492R [13]. The gene was sequenced in triplicate using Sanger

technology (Eurofins, Germany). Gene sequence (1444 bp) was deposited in the

235 GenBank/EMBL/DDBJ databases under accession number MT376293.

236 The 16S rRNA sequences from strain $MO1340^{T}$ and 11 of the closest

237 *Hydrogenothermales* obtained by BLASTN search were then edited (1364 bases) in the

238 Geneious v10.2.3 program and aligned using MUSCLE program [15, 36]. Phylogenetic

reconstructions were made on the basis of evolutionary distance using neighbour-joining [63]

240 and maximum-likelihood methods with Tamura-Nei correction (respectively modelled with G

and G+I parameters) using the MEGA 7 software [38, 75]. The reliability of internal branches
was assessed using the bootstrap method with 1000 replicates [17].

Genome sequence determination of strain MO1340^T was carried out using Illumina 243 HiSeq technology (Eurofins Scientific, Germany). Genome assembly was performed using 244 245 spades v.3.13.0 ([3]). Completion and contamination were assessed using CheckM [56] and 246 quality of the genome sequence was checked using the Quality Assessment Tool for Genome 247 Assemblies (QUAST) [30]. Genome was annotated using PGAP (Prokaryotic Genome 248 Annotation Pipeline) [76]. Average Nucleotide Identity (ANI) values were calculated with the 249 OrthoANIu algorithm [87] and can accurately replace DNA-DNA hybridization values for 250 strains for which genome sequences are available [24]. 251 252 **Results and discussion** Morphology and physiology of strain MO1340^T 253 Strain MO1340^T morphological, physiological, genomic and metabolic features and 254 255 those of its closest relatives within the phylum Aquificota, family Hydrogenothermaceae are presented in Table 1. Cells of strain MO1340^T were Gram-negative short rods from 1.4 - 2.2 256 257 $(SD=0.42 \ \mu m, n = 33) \ \mu m$ length and from 0.5 - 0.8 μm width $(SD=0.13 \ \mu m, n = 33)$, and 258 were in the range of cell sizes reported for the other *Persephonella* species (Table 1). Irregular 259 cocci were frequently observed at temperatures close to the limits for growth. TEM

260 observation showed that one-third of the cells had a single polar flagellum (Fig. 1). However,

261 motility was rarely observed, in most cases for dividing cells and in medium containing

262 elemental sulphur. On solid media, round and orange colonies (1 mm in diameter) were

263 observed after 1 to 3 days whereas cultures in liquid media were colourless.

Growth was observed for strain MO1340^T at NaCl concentrations ranging from 1.5 to 3.5 % (w/v) with an optimum at 2.5 % and at pH between 5.3 and 7.0 with an optimum

| 266 | around 6.5, suggesting these microorganisms are mostly adapted to habitats where the |
|-----|--|
| 267 | hydrothermal fluid mixes with the seawater. Although strain MO1340 ^T was characterized for |
| 268 | these parameters in presence of thiosulphate and dioxygen, optimal NaCl concentration and |
| 269 | pH for growth were similar to those obtained for all of Persephonella species grown with |
| 270 | hydrogen and nitrate (i.e. NaCl : 2.5%; pH : 6.0 to 7.2; Table 1). However, as detailed in the |
| 271 | last section "Physiological and metabolic adaptation to a contrasted environment", the |
| 272 | optimal growth temperature (i.e. 65 to 75 $^{\circ}$ C) and dioxygen dissolved concentration (i.e. 11.4 |
| 273 | to 119.7 μ M) were substrate-dependent. |
| 274 | |
| 275 | Insights into the metabolism of Persephonella strain $MO1340^{T}$ |
| 276 | Like other <i>Persephonella</i> isolates, strain MO1340 ^T was able to grow |
| 277 | chemolithoautotrophically on various inorganic catabolic substrates. This includes oxidation |
| 278 | of reduced-sulphur compounds, hydrogen-oxidation, aerobic respiration or nitrate-reduction |
| 279 | [25, 50]. The isolate grew in microaerphilic conditions with thiosulphate (10 mM), elemental |
| 280 | sulphur (1 % w/v) or hydrogen (dissolved concentration 820.7 μM). Interestingly, the optimal |
| 281 | oxygen concentrations varied according to electron donor and acceptor (optimum dissolved |
| 282 | concentrations from 11.4 to 119.7 μ M at 70 °C). Strain MO1340 ^T was also able to grow |
| 283 | anaerobically with hydrogen as electron donor (dissolved concentration 820.7 $\mu M)$ using |
| 284 | nitrate (8 mM) as final electron acceptor. In order to compare strain MO1340 ^T to its closest |
| 285 | relative, growth requirements were also tested in the same conditions using P. hydrogeniphila |
| 286 | 29W ^T (DSM 15103). Although <i>P. hydrogeniphila</i> 29W ^T was initially described as only |
| 287 | capable of using hydrogen as electron donor [50], growth kinetic showed it could also oxidise |
| 288 | thiosulphate to sulphate with oxygen as electron acceptor (Fig. SM 1, Table 2), which is |
| 289 | congruent with the presence in its genome of core genes soxABXYZ (GenBank accession |
| 290 | number: PRJEB22457). Moreover, no aerobic or anaerobic growth was observed for |

Persephonella strain *MO1340^T* in the presence of all tested organic carbon sources (yeast
extract, peptone, glucose, sucrose, fructose, lactose, or sorbitol) even in presence of carbon
dioxide. In the absence of vitamins, growth was not inhibited, even after five subcultures,
showing strain MO1340^T is a strict autotroph using only carbon dioxide as sole carbon source
as defined by Srinivasan *et al.* [71].

Substrates consumption and metabolites production by strain MO1340^T were 296 297 measured in order to establish stoichiometry of each catabolic reaction and infer the 298 associated standard Gibbs energy. Measured stoichiometry were consistent with the reactions 299 described by Amend et al. [2] (Table 2), excepted for aerobic elemental sulphur oxidation where discrepancies were observed. We hypothesize that it might result from unquantified 300 301 intracellular metabolic intermediates accumulation. For anaerobic hydrogen-depended denitrification, strain MO1340^T realised a complete reduction of nitrate to nitrogen without 302 accumulation of nitrite, nitrous oxide or ammonium which is consistent with previous studies 303 on *P. hydrogeniphila* 29W^T [50]. Moreover, in order to accurately determine the reaction 304 305 involved in aerobic hydrogen oxidation, as the end-product of aerobic hydrogen oxidation can 306 either be water $(2 H_2 + O_2 \rightarrow 2 H_2O)$ or hydrogen peroxide $(H_2 + O_2 \rightarrow H_2O_2)$ [78], hydrogen 307 peroxide production was assessed. As no production of hydrogen peroxide was detected, strain MO1340^T can probably perform the Knallgas reaction as commonly described for other 308 309 Aquificota [27, 28]. Thermodynamic calculations based on the geochemical conditions 310 measured in situ at Lucky Strike vent field showed that energy provided by this reaction could 311 reach - 280 kJ/mol (Table 2), suggesting aerobic hydrogen oxidation could be a significant 312 metabolism in hydrothermal ecosystems [44, 66]. 313 *Physiological and metabolic adaptation to a contrasted environment*

314 Growth rate measurements showed that strain MO1340^T exhibited different responses 315 to temperature depending on the catabolic reactions involved. Optimal growth temperatures

| 316 | were respectively 65, 70, 70-75 and 75 °C for the following redox couples: S^0/O_2 , $S_2O_3^2/O_2$, |
|-----|--|
| 317 | H_2/O_2 , H_2/NO_3^- . All curves for the determination of optimal growth temperature presented a |
| 318 | plateau between 5 and 15 °C below the optimal growth temperature (Fig. 2). This curve shape |
| 319 | was also previously observed for <i>P. hydrogeniphila</i> 29W ^T (supplementary data in ref. [50]) |
| 320 | that is the closest relative of strain MO1340 ^T . Arrhenius plots obtained from data of optimal |
| 321 | growth temperature experiments on strain MO1340 ^T also revealed two distinct slopes at sub- |
| 322 | optimum temperatures separated by a "critical temperature" instead of a single linear slope |
| 323 | (Fig. SM 2). Various microorganisms exhibiting growth that does not meet the square root |
| 324 | equation [57] have also been described in studies of mesophilic and thermophilic and |
| 325 | psychrophilic Bacteria [26, 48, 53]. Wiegel et al. suggested that these broken Arrhenius plots |
| 326 | could be the consequence of the expression of different sets of key enzymes capable of |
| 327 | undergoing conformational changes at different temperatures [83-85]. Moreover, the effect of |
| 328 | temperature on the specific growth rate fitted different bimodal Gaussian patterns ($R^2 = 0.96$ |
| 329 | to 1) depending on the catabolic pathway (Fig. 2). Each bimodal pair could be grouped |
| 330 | according to the composition of the electron donor involved (i.e. sulphur or hydrogen; Fig. 2). |
| 331 | For sulphur based electron donors (i.e. elemental sulphur and thiosulphate) both peaks were |
| 332 | comprised between 52.7 - 56.3 and 67.4 - 68.8 °C, whereas for hydrogen both peaks were |
| 333 | comprised between 59.8 - 63.7 and 74.1 - 74.2 (table 2 and Fig. 2), suggesting that the \approx 7 °C |
| 334 | shift depending on the electron donor could be the consequence of a different temperature |
| 335 | range for the enzymes involved in each catabolic pathways (Sox enzymes or hydrogenases). |
| 336 | For example, it was shown for Hydrogenobacter thermophilus, another member of phylum |
| 337 | Aquificota, but in the order Aquificales, which presents physiological characteristics close to |
| 338 | those of strain MO1340 ^T [35], that the Sox enzymes implied in aerobic thiosulphate oxidation |
| 339 | exhibited an optimal activity at lower temperature than the nitrite reductase (respectively 60 |
| 340 | and 70 - 75 °C) [64, 73]. |

Oxygen requirements of strain MO1340^T also depended on the electron donor used. 341 342 The optimal dissolved oxygen concentration was $11.4 \,\mu M (0.7 \,\text{mM in gas phase})$ for 343 hydrogen oxidation, and was comparable to other Persephonella species (0.73 to 1.21 mM in 344 gas phase) in the same culture conditions [25, 50]. However this value was 5 to 11 fold higher 345 in the presence of thiosulphate or elemental sulphur, respectively 59.8 and 119.7 μ M (3.68) 346 and 7.36 mM in gas phase). Comparison of growth parameters of each catabolic reaction performed by strain MO1340^T showed that optimal growth temperature and optimal oxygen 347 348 concentration were inversely correlated (table 2 and Fig. 3). For example, oxidation of 349 sulphur species such as thiosulphate and elemental sulphur allowed growth at higher oxygen concentration and at lower temperature (respectively 59.8 and 119.7 µM dissolved oxygen 350 351 and 65/70 °C) whereas hydrogen oxidation coupled to oxygen reduction was optimal with 352 only 11.4 µM oxygen at 70-75 °C. Anaerobic hydrogen oxidation coupled to nitrate reduction 353 was optimal at a higher temperature (75 °C) compared to microaerophilic metabolisms. This 354 inverse relationship between temperature and oxygen concentration was congruent with the 355 geochemical gradients observed in situ where oxygen depletion occurs as temperature 356 increases along the mixing gradient between the cold deep seawater and the hot hydrothermal 357 fluid that is expelled at the seafloor (Fig. 3).

358 In order to estimate the energy provided by catabolic reactions performed by strain 359 MO1340^T in natural settings, thermodynamic calculations were taken into consideration. We 360 used in situ data collected along the mixing between seawater and the hydrothermal fluid 361 from Lucky Strike, the site from which the strain was isolated. When data were not available, 362 we used a numerical model to predict dilution of the substrates considered along the mixing 363 gradient. The observed and predicted thermal and chemical gradients in the mixing zone were 364 used as proxies to determine the geochemical settings of hydrothermal chimneys, in order to 365 calculate the Gibbs free energy under *in situ* conditions (Δ Gr) for each catabolic reaction of

strain MO1340^T at its optimal temperature (Table 2). Elemental sulphur and thiosulphate 366 367 oxidation reactions were the most energetic (respectively -517 and -731 kJ/mol) compared to 368 aerobic and anaerobic hydrogen oxidation reactions (respectively - 280 and - 271 kJ/mol). 369 Comparison of the Δ Gr values over a large temperature range from 5 to 153 °C (Fig. SM 3) 370 showed that energy gained from elemental sulphur and thiosulphate oxidation decreased as 371 the temperature increased due to oxygen depletion, in contrast to aerobic and anaerobic 372 hydrogen oxidation, which was more energetic at higher temperature. This is consistent with optimal temperature measurements of strain MO1340^T that showed a faster growth for 373 374 oxidation of sulphur compounds at lower temperature than for aerobic and anaerobic 375 hydrogen oxidation. Moreover, for aerobic hydrogen oxidation, the optimal Δ Gr value was 376 reached at temperature above 72 °C (-280 kJ/mol, Fig. SM 3), which was congruent with the experimental results on strain MO1340^T that presented an optimal growth temperature at 70-377 378 75 °C when grown on these substrates. However, the calculated Gibbs free energies in *in situ* 379 conditions do not fully explain this phenomenon but rather provide tendencies. Several 380 reasons may explain slight discrepancies. First, the temperatures and substrate concentrations 381 used in the ΔG_r calculations are predicted from the mixing gradient between the end-member 382 hydrothermal fluid expulsed at the seafloor and the surrounding seawater, i.e. within an open 383 media. This might not exactly reproduce the conditions prevailing in a hydrothermal chimney, 384 where chemical and thermal gradient are squeezed due to the structure and porosity of the 385 chimneys [11]. For species derived from the simple dilution model (elemental sulphur, 386 hydrogen and nitrogen), biotic and abiotic reactions may occur early along the mixing 387 gradient including precipitation and solubilisation processes further altering the distribution of 388 electron donors and acceptors along dilution [40]. However, good convergence between 389 theoretical and *in situ* sulphate distribution (Fig. SM 4) gives us confidence that model 390 outputs are sufficient to set a reliable baseline for thermodynamic calculations. More complex

thermodynamic modelling including anabolic and dissipation energy requirements and effect
of substrate availability would be necessary to fully resolve microbial growth rates close to
incubation observations.

394 The physical and chemical conditions along the hydrothermal gradient shape the 395 taxonomic and functional distribution of microbial communities in hydrothermal vents [41, 396 45, 46]. For example, aerobic mesophiles are frequently observed in the periphery of 397 chimneys whereas anaerobic hyperthermophiles are present in inner parts [10, 74], with some 398 exceptions depending on the structure and porosity of chimneys [52]. Unlike strict aerobes or 399 anaerobes that could only colonize specific microhabitats in hydrothermal chimneys, metabolically versatile Aquificota strain MO1340^T could settle in contrasted parts of 400 401 chimneys due to its ability to use different electron acceptors (i.e. oxygen or nitrate). This 402 capacity to colonize numerous microhabitats in hydrothermal ecosystems was confirmed by 403 microbial diversity analysis using high-throughput sequencing that showed the presence of 404 phylotypes related to strain MO1340 in chimneys and associated fluids from the gradient at 405 five sites of the Lucky Strike vent field (unpublished results). As for Aquificota, 406 *Campylobacterota* (previously included in the former phylum *Epsilonbacteraeota*) also 407 exhibit similar catabolisms such as oxidation of reduced sulphur compounds and hydrogen 408 with both oxygen and nitrate [5]. Hence, these versatile chemolithoautotrophic microbial 409 communities thrive in a wide range of habitats along the hydrothermal gradient and therefore 410 probably drive a significant fraction of the primary production at deep-sea hydrothermal 411 systems [5, 49, 60, 68]. Moreover, it has also been suggested that a specific distribution 412 within these different taxonomic groups could also occur along the hydrothermal fluid 413 gradient. For example, within Campylobacterota, Campylobacterales seem to thrive at lower 414 temperatures (i.e. from 20 °C to 40 °C) whereas Nautiliales could colonise habitats between 415 40 °C to 60 °C [5, 20]. Aquificota could dominate niches exceeding 60 °C as they also show a 416 wide catabolic and temperature adaptation that follows the hydrothermal gradient as for strain

417 MO1340^T. This metabolic versatility and their broad temperature range could therefore

418 provide an ecological advantage in colonizing such a dynamic part of the ecosystem

419 compared to other chemoautotrophic microorganisms (Fig. 3). More comprehensive studies

420 should be conducted using combination of physiological, genomic, and ecological approaches

421 focusing this strain and others members of phylum *Aquificota* and *Epsilonbacteraeota*.

422

Genomic features and phylogenetic position

423 The draft genome of strain MO1340^T (5 contigs > 500 bp, coverage 840 x) has been

424 deposited in GenBank/EMBL/DDBJ under the accession number JAACYA000000000. The

425 genome characteristics such as size (1.892.283 bp, 99.59 % completion, 1.68 %

426 contamination) average DNA G + C content (37.07 mol%) and number of coding sequences

427 (2028 CDS) were comparable with those of previously sequenced genomes of *Persephonella*

428 strains (Table 1). ANI values with *P. hydrogeniphila* 29W^T and *P. marina* EX-H1^T

429 (respectively 75.66 % and 72.97 %) were well below the threshold (95-96 %) for

430 differentiating species [7].

Analysis of draft genome of strain MO1340^T also provided details on the metabolic 431 432 pathways involved in sulphur and hydrogen oxidation, nitrogen and oxygen reduction, and in 433 carbon fixation. Thiosulphate oxidation may be performed by the sox pathway [21, 22] as all 434 core genes *soxABXYZ* were present, excepted for *soxCD* genes that were not detected. 435 Although it was shown that organisms that lack *soxCD* genes are often unable to completely oxidise thiosulphate to sulphate [23, 67], strain MO1340^T produced sulphate in stoichiometric 436 437 quantities to the amount of thiosulphate added, suggesting an alternative pathway that could 438 complete oxidation to sulphate. Numerous genes encoding hydrogenases were also detected, 439 including [Ni-Fe]-Hydrogenases-genes, which is consistent with the ability of the strain to 440 oxidise hydrogen. More than 30 genes encoding for cytochromes (bc, c, c1, c3, cbb3 and d -

441 type) were detected and were probably specific to the strain microaerophilic lifestyle. In 442 anaerobic conditions, a complete reduction of nitrate was achieved as genes encoding nitrate 443 reductase (NapAGH), nitrite reductase (NirS), nitrite oxide reductase (Nor) and nitrous oxide 444 reductase (NosZD) were present. Autotrophic carbon dioxide fixation is performed using the 445 "A-type" reductive tricarboxylic acid (rTCA) cycle, like others members of genus 446 Persephonella and family Hydrogenothermaceae [32, 59]. As for the Calvin-Benson-447 Bassham cycle, the rTCA cycle is one of the dominant carbon fixation pathways in 448 hydrothermal vents [33, 43], although it seems more adapted to high-temperature and low oxygen environments, which is congruent with the physiology of strain MO1340^T [4, 33, 49]. 449 450 In dynamic hydrothermal ecosystems, the flagella-mediated motility and chemotaxis 451 are essential for microorganisms to respond to variation of environmental conditions (e.g. pH, 452 temperature) and to find substrates for growth (for e.g. [86]). Microscopic observations 453 showed motile cells were mainly observed in medium containing elemental sulphur, 454 suggesting a close relationship between flagella and sulphur-oxidizing systems as previously 455 observed for Acidithiobacillus spp. [86]. This link was confirmed by the presence in the genome of strain MO1340^T of several genes implied in motility or chemotaxis (*FlgGAHIJBC*, 456 457 FlhAFB, FliGMNEFR) that clustered with genes encoding transcriptional regulators, such as 458 sigma factors, or genes responsible for chemotaxis (CheZ, CheA, CheW and CheV). 459 Regulation of motility and chemotaxis by sulphur may therefore also contribute to surface 460 colonisation of microniches in hydrothermal chimneys. 461 Based on phylogenetic analyses of the 16 rRNA gene sequence (Fig. 4), strain 462 MO1340^T belongs to the genus *Persephonella*, family *Hydrogenothermaceae* order *Hydrogenothermales*. The closest described species are *P. hydrogeniphila* 29W^T, *P. marina* 463 EX-H1^T and *P. guaymasensis* EX-H2^T with respectively 95.5 %, 93.8 % and 94.0 % 16S 464 465 rRNA gene sequence similarity. These values are below the previously published cut-off

- 466 threshold of 98.7 % for species delineation, which is congruent with ANI values to other
- 467 species in the genus [7]. AAI values to other Persephonella spp. fell into the range of
- 468 interspecies AAI values confirming its novelty in the genus. On the basis of its physiological,
- 469 metabolic and genomic characteristics, we propose that strain MO1340^T (=UBOCC-M-3359
- 470 = JCM 34026) represents a novel species within the genus *Persephonella*, for which we
- 471 suggest the name *Persephonella atlantica* sp. nov, as a reference for being isolated from a
- 472 hydrothermal vent field on the Mid-Atlantic Ridge.
- 473

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481 Authors contribution

- 482 David François: Investigation, Writing Original Draft. Anne Godfroy: Supervision,
- 483 Samples collection. Clémentine Mathien: Investigation. Johanne Aubé: Genome assembly.
- 484 **Cécile Cathalot**: Thermodynamic modelling. **Françoise Lesongeur**: Samples collection.
- 485 Stéphane l'Haridon: Investigation, Methodology. Xavier Philippon: Investigation. Erwan
- 486 Roussel: Conceptualization, Investigation. All authors: Writing Review



491 Figure 1. Transmission electron micrograph of a cell of *Persephonella atlantica* strain

492 MO1340^T showing the polar flagellum. Scale bar represents $0.5 \,\mu$ m.





502



- 504 Figure 3. Schematic comparison of optimal temperatures and O₂ concentrations for
- 505 *Persephonella atlantica* strain MO1340^T depending on catabolic substrates and *in situ* thermal
- 506 and chemical gradients observed during the mixing of hydrothermal fluid and seawater.
- 507 Concentration of chemicals species correspond to the maximal concentrations measured in
- 508 situ or calculated for the hydrothermal physico-chemical gradient between 5 and 153 °C
- 509 (details in section *Thermodynamic calculations and Arrhenius parameters*).
- 510



- 511 0.020
- 512 Figure 4. Phylogenetic tree of members of order *Aquificales* based on 16S rRNA gene
- 513 sequences (1364 bp) showing the position of *Persephonella atlantica* strain MO1340^T among
- 514 related species. Trees reconstructed using the neighbour-joining (NJ) and maximum-
- 515 likelihood (ML) methods displayed the same topology. Bootstrap values (shown in the order

- NJ/ML) are given to the left of each node. Bar represents 0.02 changes per nucleotide position. Numbers in parentheses are GenBank/EMBL/DDBJ accession numbers.
- 516 517 518

519 Table 1. Comparison of some characteristics of strain MO1340^T with other closest marine

520 strains within the family *Hydrogenothermaceae*. Strain MO1340^T (this study); *Persephonella*

521 *hydrogeniphila* 29W^T [50]; *P. marina* EX-H1^T [25]; *P. guaymasensis* EX-H2^T [25];

522 *Hydrogenothermus marinus* VM1^T [72]; +, positive; -, negative; *, only with O_2 as electron

523 acceptor; **, shown in this study; ^a, culture grown with S^0/O_2 ; ^b, culture grown with $S_2O_3^2/O_2$; ^c,

524 culture grown with H_2/O_2 ; ^d, culture grown with H_2/NO_3 ; \cong Inconsistent growth; ND, not

determined. Oxygen concentration presented was calculated in gas phase. DNA G+C content
 was calculated from genome analysis.

| Characteristics | Strain MO1340 | P. hydrogeniphila 29W [™] | P. marina EX-H1 [⊤] | P. guaymasensis EX-H2 [⊤] | Hydrogenother mus marinus VM1 [™] |
|-----------------------------------|--|--|--|--|--|
| Cell morphology | rods | rods - cocci | rods | rods | rods |
| Cell size (length/width μm) | 1.4-2.2 / 0.5- 0.8 | 2.3-2.7 / 0.9-1 | 2-4 / 0.3- 0.4 | 2-4 / 0.3-0.4 | 2-4 / 1-1.5 |
| Motility | + | + | + | + | + |
| Temperature range | 50-75ª ; 45-80 ^{bd} ; 55-80 ^c | 50-72.5 ^d | 55-90 ^d | 60-80 ^d | 65° |
| Temperature optimum (°C) | 65ª ; 70 ^b ; 70- 75 ^c ; 75 ^d | 70 ^d | 70 ^d | 75 ^d | 45-80 ^c |
| pH optimum | 6.5 ^b | 7.2 ^d | 6 ^d | 6 ^d | 5-7° |
| pH range | 5.30 - 7 ^b | 5.5-7.6 ^d | 4.7-7.5 ^d | 4.7-7.5 ^d | 5-7 ^c |
| NaCl optimum (% w/v) | 2.5 ^b | 2.5 ^d | 2.5 ^d | 2.5 ^d | 2-3 ^c |
| NaCl range (% w/v) | 1.5-3.5 ^b | 1.5-5.0 ^d | 1.0-4.5 ^d | 1.0-4.5 ^d | 0.5-6 ^c |
| O2 optimum (mM) | 7.36 ^a ; 3.68 ^b ; 0.7 ^c | 0.63-0.84 ^c | 0.97-1.45 ^c | 0.97-1.45 ^c | 0.69-1.38 ^c |
| O2 range (mM) | 0.52-> 11ª ; 0,52-10.5 ^b ; ND-3.5 ^c | 0.10-1.05° | ND-< 4.53 ^c | ND-< 5.32 ^c | 0.34-5.53 ^c |
| Electron donors | H ₂ , S ₂ O ₃ ²⁻ *, S ⁰ * | H ₂ , S ₂ O ₃ ^{2-**} | H ₂ , S ₂ O ₃ ^{2-*} , S ^{0*} | H ₂ , S ₂ O ₃ ²⁻ *, S ^{0*} | H ₂ |
| Electron acceptors | NO ₃ ⁻ , O ₂ | NO ₃ ⁻ , O ₂ | NO ₃ ⁻ , O ₂ , S ⁰ , SO ₄ ²⁻ \aleph , acetate \aleph | NO ₃ -, O ₂ | O ₂ |
| Genome accession number | JAACYA000000 000 | NZ_OBE100000 000 | NC_012440 | - | NZ_REFO00000 000 |
| G + C content (mol%) | 37.1 | 35.1 | 37.1 | - | 27.7 |
| Size (Mb) | 1.89 | 2.0 | 1.98 | - | 1.60 |
| No. of contigs | 5 | 19 | 2 | - | 19 |
| No. of ORFs | 2028 | 2130 | 2083 | - | 1713 |

⁵²⁷

528

530 Table 2. Summary of catabolic reactions performed by *Persephonella atlantica* strain

531 MO1340^T and physiological parameters according to substrates couples. \bowtie Optimal

temperature corresponding to the two peaks observed by bimodal Gauss fitting of growth

533 curves reported in figure 3. Putative *in situ* Δ Gr values at 72 °C were calculated at optimal

534 growth temperature of the reactions, using the high concentration range from Rommeveaux *et*

535 *al* [61].

536

| Reaction | Optimal growth temperature (range) °C | 1st and 2nd peak from bimodal fitted curves (°C) ¤ | Optimal O ₂ concentration in gas phase (range) mM | Optimal dissolved O ₂ concentration (range) µM | Q ₁₀ (range of linearity on Arrhenius plot, °C) | ∆Gr (kJ/mol) |
|---|--|---|---|--|---|-----------------|
| $\begin{array}{c} \mathrm{S}^{0} + 1.5 \ \mathrm{O}_{2} + \mathrm{H}_{2}\mathrm{O} \rightarrow \\ \mathrm{SO}_{4}{}^{2\text{-}} + 2 \ \mathrm{H}^{+} \end{array}$ | 65 (55-80) | 56.3- 67.4 | 7.36 (0.52-> 11) | 119.7 (8.5-> 179.5) | 1,53 (55-65) | -517 |
| $\begin{split} S_2 O_3^{2\text{-}} &+ 2 \ O_2 + H_2 O \\ &\rightarrow 2 \ SO_4^{2\text{-}} + 2 \ H^+ \end{split}$ | 70 (45-80) | 52.7- 68.8 | 3.68 (0.52-10.5) | 59.8 (8.5-170.9) | 1,18 (50-70) | -731 |
| $2 \text{ H}_2 + \text{O}_2 + \rightarrow 2 \text{ H}_2\text{O}$ | 70-75 (55-80) | 63.7- 74.1 | 0.7 (ND-< 3.5) | 11.4 (ND-< 57) | 1,3 (60-75) | -280 |
| $\begin{array}{c} 2.5 \ H_2 + NO_3^- + H^+ \rightarrow \\ 0.5 \ N_2 + 3 \ H_2O \end{array}$ | 75 (45-80) | 59.8- 74.2 | ND | ND | 1,33 (55-75) | -271 |

| 538 | Table 3. Description | of Persephonella | atlantica sp. nov. |
|-----|----------------------|------------------|--------------------|
|-----|----------------------|------------------|--------------------|

| Genus name | Persephonella |
|---|---|
| Species name | Persephonella atlantica |
| Specific epithet | atlantica |
| Species status | sp. nov. |
| Species etymology | at.lan'ti.ca. L. fem. adj. atlantica for being isolated from the Atlantic Ocean |
| Description of the new taxon and diagnostic traits | Gram-negative cells, motile short rods with a mean length of 1.4-2.2 μ m and a width of 0.5-0.8 μ m. Colonies are orange, round and regular with a diameter of 1 mm. Strict chemolithoautotroph growing in microaerophilic or anaerobic conditions. No heterotophic or mixotrophic growth was observed. Vitamins or organic chelators are not necessary for growth. Uses hydrogen, thiosulphate or elemental sulphur with oxygen as electron acceptor. Sulphate is produced from oxidation of sulphur compounds. Hydrogen can also be coupled to nitrate reduction, producing nitrogen. Growth was observed for strain MO1340 ^T at NaCl concentrations ranging from 1.5 to 4.5 % (w/v) with an optimum at 2.5 % and at pH between 5.3 and 7.0 with an optimum around 6.5. Optimal growth temperature is respectively 65, 70, 70-75 and 75 °C, and optimal dissolved oxygen concentration is 119.7, 59.8, 11.4 and <0.2 μ M for the following redox couples: S ⁰ /O ₂ , S ₂ O ₃ ²⁻ /O ₂ , H ₂ /O ₂ , H ₂ /NO ₃ ⁻ . |
| Country of origin | EEZ Portugal |
| Region of origin | Lucky Strike hydrothermal field |
| Date of isolation (dd/mm/yyyy) | 29/01/2018 |
| Source of isolation | Black smoker chimney |
| Sampling date (dd/mm/yyyy) | 12/08/2008 |
| Latitude (xx°xx′xx″N/S) | 37° 17' 20.26" N |
| Longitude (xx°xx′xx″E/W) | 32° 16' 32.05" W |
| 16S rRNA gene accession nr. | MT376293 |
| Genome accession number | JAACYA00000000 |
| Genome status | Incomplete (5 contigs > 500 bp) |
| Genome size | 1.892.283 bp |
| GC mol% | 37.07 |
| Number of strains in study | 1 |
| Information related to the Nagoya Protocol | MOMAR 2008 oceanographic cruise. Nota verbal n°2406 June 02 2008 from ministerio dos négocios estrangeiros Portugal |
| Designation of the Type Strain | MO1340 ^T |
| Strain Collection Numbers | UBOCC-M-3359 = JCM 34026 |

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