

Methanonatronarchaeum thermophilum gen. Nov., sp. nov. and 'Candidatus methanohalarchaeum thermophilum', extremely halo(natrono)philic methyl-reducing methanogens from hypersaline lakes comprising a new euryarchaeal class Methanonatronarchaeia classis nov.

Sorokin, Dmitry Y.; Merkel, Alexander Y.; Abbas, Ben; Makarova, Kira S.; Rijpstra, W. Irene C.; Koenen, M.; Sinninghe Damsté, Jaap S.; Galinski, Erwin A.; Koonin, Eugene V.; Van Loosdrecht, Mark C.M.

DOI

[10.1099/ijsem.0.002810](https://doi.org/10.1099/ijsem.0.002810)

Publication date

2018

Document Version

Accepted author manuscript

Published in

International Journal of Systematic and Evolutionary Microbiology

Citation (APA)

Sorokin, D. Y., Merkel, A. Y., Abbas, B., Makarova, K. S., Rijpstra, W. I. C., Koenen, M., Sinninghe Damsté, J. S., Galinski, E. A., Koonin, E. V., & Van Loosdrecht, M. C. M. (2018). Methanonatronarchaeum thermophilum gen. Nov., sp. nov. and 'Candidatus methanohalarchaeum thermophilum', extremely halo(natrono)philic methyl-reducing methanogens from hypersaline lakes comprising a new euryarchaeal class Methanonatronarchaeia classis nov. *International Journal of Systematic and Evolutionary Microbiology*, 68(7), 2199-2208. Article 002810. <https://doi.org/10.1099/ijsem.0.002810>

Important note

To cite this publication, please use the final published version (if applicable).
Please check the document version above.

Copyright

Other than for strictly personal use, it is not permitted to download, forward or distribute the text or part of it, without the consent of the author(s) and/or copyright holder(s), unless the work is under an open content license such as Creative Commons.

Takedown policy

Please contact us and provide details if you believe this document breaches copyrights.
We will remove access to the work immediately and investigate your claim.

International Journal of Systematic and Evolutionary Microbiology
Methanonatronarchaeum thermophilum gen. nov., sp. nov, and 'Candidatus
Methanohalarchaeum thermophilum', extremely halo(natrono)philic methyl-reducing
methanogens from hypersaline lakes comprise a new euryarchaeal class
Methanonatronarchaeia classis nov.

--Manuscript Draft--

Manuscript Number:	
Full Title:	Methanonatronarchaeum thermophilum gen. nov., sp. nov, and 'Candidatus Methanohalarchaeum thermophilum', extremely halo(natrono)philic methyl-reducing methanogens from hypersaline lakes comprise a new euryarchaeal class Methanonatronarchaeia classis nov.
Article Type:	Taxonomic Description
Section/Category:	New taxa - Archaea
Keywords:	methanogenesis methyl-reducing hypersaline salt lakes soda lakes Methanonatronarchaeia
Corresponding Author:	Dimitry Y Sorokin, Ph.D., Dr.Sci. Winogradsky Institute of Microbiology, Research Centre of Biotechnology RAS Moscow, NA RUSSIAN FEDERATION
First Author:	Dimitry Y Sorokin, Ph.D., Dr.Sci.
Order of Authors:	Dimitry Y Sorokin, Ph.D., Dr.Sci. Alexander Y Merkel, PhD Ben Abbas, MSc Kira S Makarova, PhD Irene W.C. Rijpstra Michel Koenen Jaap S. Sinninghe Damsté, PhD Erwin A. Galinski, PhD Eugene V. Koonin, PhD Mark C.M. van Loosdrecht, Eng.,PhD
Manuscript Region of Origin:	RUSSIAN FEDERATION
Abstract:	Methanogenic enrichments from hypersaline lakes at moderate thermophilic conditions resulted in cultivation of an unknown deep lineage of euryarchaeota related to the class Halobacteria. Both soda and salt lake isolates belong to methyl-reducing methanogens that utilize C1 methylated compounds as electron acceptor and H ₂ or formate as electron donor. They are extreme halophiles, growing optimally at 4 M total Na ⁺ and represent the first example of methanogens employing the "salt-in" osmoprotection mechanism. The salt lake subgroup is neutrophilic, whereas the soda lake isolates are obligate alkaliphiles, with an optimum around pH 9.5. Both grow optimally at 50°C. The genetic diversity inside the two subgroups is very low, indicating that the soda and salt lake clusters consist of a single genetic species each. The phylogenetic distance between the two subgroups is in the range of distant genera, whereas the distance to other euryarchaea is below 83% identity of the 16S rRNA. These isolates and closely related environmental clones from hypersaline habitats (SA1 group) form a novel class-level clade in the phylum Euryarchaeota that is strongly supported by bootstrap analysis. On the basis of distinct phenotypic and genetic properties, the soda lake isolates are classified into a new genus and species

Methanonatronarchaeum thermophilum with the type strain AMET1T (DSM 26684T=NBRC 110805T=UNIQEM U982), and the salt lake methanogens - as a candidate genus and species 'Ca. Methanohalarchaeum thermophilum'. Together with uncultured SA1 group clones, these organisms are proposed to form a new class Methanonatronarchaeia within the phylum Euryarchaeota.

2 ***Methanonatronarchaeum thermophilum* gen. nov., sp. nov, and 'Candidatus**
3 ***Methanohalarchaeum thermophilum*', extremely halo(natrono)philic**
4 **methyl-reducing methanogens from hypersaline lakes comprise a new**
5 **euryarchaeal class *Methanonatronarchaeia* classis nov.**
6
7

8 Dimitry Y. Sorokin^{a,b*}, Alexander Y. Merkel^a, Ben Abbas^b, Kira S. Makarova^c, W. Irene C.
9 Rijpstra^d, M. Koenen^d, Jaap S. Sinninghe Damsté^{d,e}, Erwin A. Galinski^f, Eugene V. Koonin^c
10 and Mark C.M. van Loosdrecht^b
11

12 ^a*Winogradsky Institute of Microbiology, Research Centre of Biotechnology, Russian Academy of Sciences,*
13 *Moscow, Russia*

14 ^b*Department of Biotechnology, TU Delft, The Netherlands*

15 ^c*National Center for Biotechnology Information, National Library of Medicine, National Institutes of Health,*
16 *Bethesda, MD, USA*

17 ^d*Department of Marine Microbiology and Biogeochemistry, NIOZ Netherlands Institute for Sea Research, and*
18 *Utrecht University, The Netherlands*

19 ^e*Department of Earth Sciences – Geochemistry, Faculty of Geosciences, Utrecht University, Utrecht, The*
20 *Netherlands*

21 ^f*Institute of Microbiology and Biotechnology, Rheinische Friedrich-Wilhelms University, Bonn, Germany*
22
23

24 *Author for correspondence:

25 D.Y. Sorokin; Tel: (7-495)1350109, Fax: (7-495)1356530; e-mail: soroc@inmi.ru; d.sorokin@tudelft.nl
26
27
28
29
30
31

32 Running title: *Methanonatronarchaeum thermophilum* gen. nov., sp. nov, and 'Candidatus
33 *Methanohalarchaeum thermophilum*'
34
35
36

37 The genome of the type strain AMET1^T and the metagenome of HMET1^T have been
38 deposited in the GenBank under the numbers MRZU00000000 and MSDW00000000,
39 respectively. The 16S-rRNA gene sequences of the AMET strains are deposited under the
40 numbers KY449317-KY4493127.

41 Methanogenic enrichments from hypersaline lakes at moderate thermophilic conditions
42 resulted in cultivation of an unknown deep lineage of euryarchaeota related to the class
43 *Halobacteria*. Both soda and salt lake isolates belong to methyl-reducing methanogens
44 that utilize C₁ methylated compounds as electron acceptor and H₂ or formate as electron
45 donor. They are extreme halophiles, growing optimally at 4 M total Na⁺ and represent
46 the first example of methanogens employing the "salt-in" osmoprotection mechanism.
47 The salt lake subgroup is neutrophilic, whereas the soda lake isolates are obligate
48 alkaliphiles, with an optimum around pH 9.5. Both grow optimally at 50°C. The genetic
49 diversity inside the two subgroups is very low, indicating that the soda and salt lake
50 clusters consist of a single genetic species each. The phylogenetic distance between the
51 two subgroups is in the range of distant genera, whereas the distance to other
52 euryarchaea is below 83% identity of the 16S rRNA. These isolates and closely related
53 environmental clones from hypersaline habitats (SA1 group) form a novel class-level
54 clade in the phylum Euryarchaeota that is strongly supported by bootstrap analysis. On
55 the basis of distinct phenotypic and genetic properties, the soda lake isolates are
56 classified into a new genus and species *Methanonatronarchaeum thermophilum* with the
57 type strain AMET1^T (DSM 26684^T=NBRC 110805^T=UNIQEM U982), and the salt lake
58 methanogens - as a candidate genus and species 'Ca. Methanohalarchaeum
59 thermophilum'. Together with uncultured SA1 group clones, these organisms are
60 proposed to form a new class *Methanonatronarchaeia* within the phylum Euryarchaeota.

61

62

63

64

65

66 In hypersaline habitats, methylotrophic methanogenesis is usually considered to be the
67 dominant pathway [1-2]. The organisms responsible for this process are members of the order
68 *Methanosarcinales*. In neutral pH conditions, *Methanosarcinales* are represented by the high
69 salt-tolerant genera *Methanohalophilus* and *Methanohalobium* that can grow at up to 4 M
70 NaCl [2-4], and a single methylotrophic genus *Methanonatronum* has been identified that can
71 grow in hypersaline soda brines [5-8]. All these methanogens, although able to tolerate salt-
72 saturating conditions, belong to moderate halophiles that grow optimally at salinity around 2
73 M total Na⁺ and utilize the bacterial type of osmoprotection based on organic compatible
74 solutes [9-10].

75
76 Our recent exploration of methanogenic archaea in sediments of hypersaline inland lakes has
77 shown that, at elevated temperatures, a previously unknown group of extremely
78 halo(natrono)philic methanogens started to outcompete the salt-tolerant *Methanosarcinales*
79 members when formate was supplied on the top of C₁ methylated compounds as
80 methanogenic substrate. This suggested the methyl-reducing nature of the novel group [11].
81 In this hybrid methanogenic pathway, the C₁ methylated compounds are used as electron
82 acceptors only, whereas external H₂ is required as the electron donor. This pathway, until
83 recently, had been considered rare, having been characterized in only two species of
84 methanogens, *Methanosphaera stadtmanae* (*Methanobacteriales*) and *Methanomicrococcus*
85 *blatticola* (*Methanosarcinales*) [12-14]. However, virtually all recent discoveries of novel
86 deep lineages of methanogens involve methyl-reducers, including the *Thermoplasmata*
87 methanogens [15-16], the Candidate class "Methanofastidiosa" [17] and the Candidate phyla
88 "Bathyarchaeota" [18] and "Verstraetearchaeota" [19]. These findings indicate that methyl-
89 reduction has so far been overlooked as an important methanogenic pathway that might be
90 able to compete with both classical methylotrophic and lithotrophic pathways. Here, we

91 describe the phenotypic and genetic properties of the novel group of extremophilic methyl-
92 reducing methanogenic euryarchaea from hypersaline lakes which we propose as founding
93 members of a new class *Methanonatronarchaeia*.

94

95 The source of the isolates was surface layer (5-15 cm) of anaerobic sediments from
96 hypersaline salt and soda lakes from various geographical locations as shown in **Table 1**.

97 Overall, eleven pure cultures of haloalkaliphilic and three highly enriched cultures of
98 halophilic methyl-reducing methanogens were obtained at 4 M total Na⁺ and 37-60°C.

99

100 The extremely haloalkaliphilic methyl-reducing AMET isolates were enriched and further
101 purified by serial dilution using mineral base medium containing 4 M total Na⁺ (2 M Na⁺ as
102 sodium carbonates + 2 M NaCl), 5 g l⁻¹ KCl and and 1 g l⁻¹ of K₂HPO₄ at pH 9.5 (4 mM
103 NH₄Cl was added after sterilization). The extremely halophilic, neutrophilic methyl-reducing
104 HMET cultures were enriched in 4 M NaCl/5 g l⁻¹ KCl, buffered at pH 7 by K₂HPO₄-KH₂PO₄
105 (total 3 g l⁻¹) and supplemented with 0.5 g l⁻¹ of NH₄Cl. After sterilization, both types of the
106 mineral basic media were supplemented with two trace metal solutions, MgCl₂ and vitamins
107 as described previously [8]. Further additions included CoM (50 μM), yeast extract (100 mg/l)
108 and either 0.1 mM hydrotroillite (FeS x nH₂O) or heat-sterilized anaerobic sediment slurries
109 (approx. 10 cm³ l⁻¹) from either soda lakes or salt lakes. 50 mM each of MeOH and sodium
110 formate were added as substrates and 0.5 mM sodium sulfide as a reductant. The media were
111 dispensed into serum bottles (from 30 to 100 ml) at 75% volume capacity and made anoxic by
112 sterile argon flushing-evacuation. Final reduction of the media was achieved by adding by
113 syringe of a drop/10 ml of 10% dithionite solution in 1 M NaHCO₃. The colonial growth of
114 strain AMET1^T was achieved in soft agar by mixing the 4 M complete sterile liquid alkaline
115 medium and 4% agarose (0.8% final) at 50°C and pouring 20 ml portions onto plates

116 containing 0.1 ml of serially diluted, fully grown liquid culture. All manipulations were
117 performed in an anaerobic glove box. The plates were incubated for one month in closed jars
118 under Ar atmosphere with the O₂-scavenging catalyzer (Oxoid) at 45°C.

119
120 Phase contrast microphotographs were obtained using a Zeiss Axioplan Imaging 2
121 microscope (Göttingen, Germany). For the total cell electron microscopy, the cells were
122 centrifuged and resuspended in 3 M NaCl, fixed with paraformaldehyde (final concentration
123 3%, v/v) for 2 h at room temperature, then washed again with the same NaCl solutions. The
124 fixed cells were positively contrasted with 1% (w/v) uranyl acetate. For thin sectioning, the
125 cell pellets were fixed in 1% (w/v) OsO₄ containing 3.0 M NaCl for 1 week at 4°C, washed
126 and resuspended in 3 M NaCl, stained overnight with 1% (w/v) uranyl acetate, dehydrated in
127 ethanol series, and embedded in Epon resin. Thin sections were poststained with 1% (w/v)
128 lead citrate. The core membrane lipids were obtained by acid hydrolysis (5% HCl in methanol
129 by reflux for 3 h) of the freeze-dried cells and subsequent analysis by HPLC-MS for GDGTs
130 and archaeol derivatives according to [20]. Intact polar lipids were obtained by Bligh Dyer
131 extraction of freeze-dried cells and subsequent HPLC-MS analysis as described in [21]. The
132 presence of intracellular organic compatible solutes in the lyophilized cells of strain AMET1^T
133 was analyzed by HPLC and ¹H-NMR after extraction with EtOH and the intracellular
134 potassium concentration was measured using ICP-MS. The cell protein was analyzed by the
135 Lowry method after removal of cell-bound FeS by several washing with acidic 4 M NaCl
136 solution.

137
138 In total, eleven pure and three highly enriched mixed methanogenic cultures were isolated
139 from anaerobic sediments of hypersaline soda and salt lakes, respectively, at 4 M total Na⁺
140 and methyl-reducing conditions, i.e. with either methanol or trimethylamine (TMA) as the

141 electron acceptor and formate or H₂ as the electron donor (**Table 1**). In addition to the
142 extreme salinity, all but one AMET strain were enriched and isolated at elevated
143 temperatures, between 48 and 60°C. The soda lake AMET isolates were successfully purified
144 from bacterial satellites using a combination of antibiotic treatment (streptomycin +
145 vancomycin, or rifampicine 100 mg l⁻¹ each) and filtration (0.45 µm). For the salt lake HMET
146 cultures, although this procedure efficiently eliminated bacteria, a small fraction (approx. 5%)
147 of other, non-methanogenic haloarcheal cells persisted in the serial dilutions. Furthermore, the
148 growth rate and yield of the HMET cultures were extremely low compared to the AMET
149 isolates, which made their purification problematic. All cultures exhibited obligate
150 dependence on external CoM and FeS. Moreover, only three of the eleven AMET isolates
151 grew in the minimal medium with these additions, whereas the rest were dependent on the
152 presence of sterilized sediments either from soda (AMET) or salt (HMET) lakes. What
153 exactly these organisms needed from the sediments, remains unclear, although a test with
154 separated pore brines and the solid phase demonstrated that the latter was far more efficient as
155 a growth factor. To our knowledge, similar observations have been reported in only one other
156 case, for an unidentified methylotrophic methanogenic culture obtained from alkaline saline
157 Mono Lake, but that culture most probably belonged to a classical methylotroph, because it
158 grew in the presence of methanol as the only substrate [22].

159 In a single case of strain AMET1^T, which can grow in presence of FeS alone, colonial
160 growth was achieved. The colonies were disc-shaped, up to 1 mm in diameter and yellow-
161 colored. The typical cell morphologies of the AMET and HMET type strains are shown on
162 **Figs. 1** and **2**. The cells are irregular angular cocci of a characteristic small size (mean cell
163 diameter is 0.4 µm). The cells of AMET strains were motile and possess multiple archaella,
164 whereas no motility was observed for the cells in HMET cultures. Both groups have a thin,
165 monolayer cell wall covered with a thick EPS layer. In addition, invaginations of cytoplasmic

166 membrane and large electron transparent inclusions (possibly, polyhydroxyalkanoates) were
167 visible in the cells of HMET1. The cells lysed immediately upon downshift in salinity below
168 2 M Na⁺. The absence of blue autofluorescence indicated the absence of deazoflavine (F₄₂₀)
169 normally present in classical methylotrophic methanogens.

170 The analysis of organic compatible solutes in cells of strain AMET1^T (grown at 4 M
171 Na⁺, pH 9.5) gave negative results. However, intracellular cation analysis demonstrated molar
172 concentrations of K⁺. These observations indicate that the novel methanogens employ the
173 haloarchaeal type (“salt-in”) osmoprotection mechanism which has not yet been demonstrated
174 for any other halophilic methanogens.

175 The core membrane lipids of AMET1^T and HMET1^T are primarily composed of a
176 mixture of GDGT-0 and archaeol (C20-C20 diphytanylglycerol diether) (**Supplementary**
177 **Table S1**). AMET1 also contained small quantities of GDGT-1, which was not detected in
178 HMET1^T. In addition to archaeol, minor amounts of two monophytanyl glycerol ethers (2-
179 C20 MGE and 1-C20 MGE) in AMET1^T and only 1-C20 MGE in HMET1^T were detected.
180 The complete absence of extended archaeols (C20-C25 and C25-C20 DGE) in membrane
181 lipids differentiated the extremely halophilic methanogens from haloarchaea [23]. The intact
182 polar lipid compositions of the two organisms were clearly different. In the alkaliphilic
183 AMET1^T, the dominant polar lipids were phosphatidylglycerol (PG) and PG-PG with GDGT-
184 0 as the core lipid. In the halophilic strain HMET1^T, the dominant polar lipids were identified
185 as dihexose derivatives of both archaeol and GDGT-0 (**Supplementary table S1**).

186 Both AMET and HMET strains use the methyl-reducing pathway of methanogenesis,
187 whereby the C₁ methylated compounds, such as methanol, methylamines or methylated
188 sulfides are used only as electron acceptors, whereas H₂ serves as the external electron donor.
189 For the AMET strains the best electron acceptor was methanol. Methylamines, including
190 mono-, di- and trimethylamine and tetramethylammonium, can also be utilized in ammonia-

191 free media but were highly toxic at alkaline conditions and the growth was much less active.
192 The growth with dimethylsulfide demanded gradual adaptation starting from 2 mM, but after
193 several steps, the best adapted strain, AMET6-2, was able to grow in presence of up to 20 mM
194 DMS. On the other hand, although possible in principle, the utilization of methanethiol was
195 irregular and no adaptation was observed to this toxic methylated compound. The neutrophilic
196 HMET strains preferred trimethylamine as the acceptor over methanol and growth with the
197 other C₁ methyl compounds was not observed. The two groups also differed in their preferred
198 *e*-donor: while the AMET strains clearly preferred formate, the HMET strains used H₂ more
199 actively. Utilization of formate as the *e*-donor, as well as DMS as the acceptor, have not been
200 demonstrated previously for any cultured methyl-reducing methanogens.

201 A unique property of the novel methyl-reducing methanogens is their extreme
202 halophily. Both groups grew within the range of Na⁺ concentrations that, among the cultured
203 archaea, are typical only for haloarchaea, i.e. from 3 to 5 M, with an optimum at
204 approximately 4 M. This preferred range of salt concentration is compatible with the evidence
205 indicating that these organisms employ the "salt-in" strategy for osmoprotection. The AMET
206 group from soda lakes belongs to obligate alkaliphiles growing within the pH range (at 4 M
207 Na⁺ and 48°C) from 8.2 to 10.2 (optimum at 9.5-9.8). In contrast to most of the extremely
208 natronophilic bacteria isolated from hypersaline soda lakes, the new archaea depend on molar
209 concentrations of NaCl and grow optimally in a medium containing 2 M NaCl and 2 M (Na)
210 carbonates. The HMET strains were typical neutrophiles with a pH range for growth from 6.5
211 to 8. Furthermore, both groups preferred elevated temperatures for growth despite being
212 isolated from moderate habitats. They grew optimally at 50°C and some of the strains
213 tolerated up to 60°C.

214 The Maximum Likelihood phylogenetic tree of 16S rRNA was constructed using
215 PhyML 3.0 with the Smart Model Selection [24], the SPR (Subtree Pruning and Regrafting)

216 type of tree improvement [25] and the aLRT (Approximate likelihood-ratio test) for branch
 217 support [26]. Only nearly complete sequences of 16S rRNA genes from the SILVA database
 218 [27] were included in the calculation. The results show that the AMET and HMET groups
 219 form two compact clades, with a maximum distance inside the groups of 1.5%. The distance
 220 between the two groups was about 10%, indicating that they represent two distinct genera.
 221 However, no close relatives of these organisms were identified among the cultivated members
 222 of Euryarchaeota, whereas among uncultured archaeal clones, the novel methanogens were
 223 clearly related to the SA1 group detected in various hypersaline habitats [28-30]. Further
 224 phylogenetic reconstruction [11] showed that the closest relatives of the AMET-HMET group
 225 in Euryarchaeota were haloarchaea of the class *Halobacteria* (**Fig. 3**) which, again, is
 226 compatible with the extreme halophily and the likely "salt-in" osmotic strategy of the novel
 227 methanogens.

228

229 Overall, on the basis of phylogenetic analysis and unique phenotypic properties, the novel
 230 moderately thermophilic and extremely halo(alkali)philic methyl-reducing methanogens from
 231 hypersaline lakes are proposed to form a new euryarchaeal class *Methanonatronarchaeia*
 232 including the alkaliphilic AMET isolates from soda lakes as a new genus and species
 233 *Methanonatronarchaeum thermophilum*, and a candidate genus and species '*Ca.*
 234 *Methanohalarchaeum thermophilum*' from salt lakes.

235

236 **DESCRIPTION OF METHANONATRONARCHAEUM GEN. NOV.**

237 Metha.no.na.tron.ar.chae'um. N.L. n. *methanum* [from French n. *méth(yle)* and chemical
 238 suffix *-ane*], methane; N.L. pref. *methano-*, pertaining to methane; N. Gr. n. *natron*,
 239 arbitrarily derived from the Arabic n. *natrun* or *natron*, soda; N. L. neut. n. *archaeum* [from
 240 Gr. adj. *archaios*, *-e*, *-on* ancient] archaeon; N. L. neut. n. *Methanonatronarchaeum* a soda-
 241 loving archaeon forming methane

242

243 Extremely halo(alkali)philic and moderately thermophilic methanogens that use the methyl-
 244 reducing pathway of methanogenesis. Utilize the "salt in" osmoprotection strategy. Found in
 245 hypersaline alkaline lakes. Member of the phylum Euryarchaeota.

246 **DESCRIPTION OF *METHANONATRONARCHAEUM THERMOPHILUM* SP. NOV.**
 247 ther.mo.phi'lum Gr. adj. *thermos*, hot; N. L. adj. *philum* (from Gr. adj. *philos* -ê -on), friend,
 248 loving; N. L. adj. *thermophilum*, thermophilic).
 249

250 The species description is based on eleven isolates. Cells are small irregular cocci, 0.4-0.5 µm
 251 in size, motile by 1-5 archaella. The cell wall is a thin monolayer covered with EPS. The cells
 252 lyze at salinity below 2 M Na⁺. Accumulate potassium as compatible solute. The F₄₂₀-
 253 dependent cell autofluorescence is absent. The colonies are yellowish, lens-shaped, up to 1
 254 mm. The core lipids are dominated by archaeol (C20-C20 DGE). Strictly anaerobic
 255 methanogens utilizing MeOH, methylamines and dimethylsulfide as electron acceptor and
 256 formate or H₂ as electron donor. Heterotrophic, can utilize yeast extract or acetate as C-
 257 source. Growth depends on external CoM, FeS/or sterilized anaerobic sediments from soda
 258 lakes. Obligately alkaliphilic with a pH range for growth from 8.2 to 10.2 (optimum at pH
 259 9.5-9.7) and extremely halo(natrono)philic, growing optimally at 4 M total Na⁺. Moderately
 260 thermophilic, with an optimum at 50°C and the upper limit for growth at 60°C. The G + C
 261 content of the genomic DNA in the type strain is 38 mol% (genome). The type strain,
 262 AMET1^T (DSM 26684=NBRC 110805=UNIQEM 982), was isolated from sediments of
 263 hypersaline soda lakes in Kulunda Steppe (Altai, Russia). The accession number of 16S
 264 rRNA gene sequence of the type strain in GenBank is KY449317.
 265

266 **DESCRIPTION OF 'CANDIDATUS METHANOHALARCHAEUM**
 267 **THERMOPHILUM'**

268 Metha.no.hal.ar.chae'um. N. L. n. *methanum* [from French n. *méth(yle)* and chemical suffix -
 269 *ane*], methane; N. L. pref. *methano-*, pertaining to methane; Gr. n. *hals halos*, sea salt, NaCl;
 270 N. L. neut. n. *archaeum* [from Gr. adj. *archaios*, -e, -on ancient] archaeon; N. L. neut.
 271 n. *Methanohalarchaeum* a salt-loving archaeon forming methane;
 272 ther.mo.phi'lum Gr. adj. *thermos*, hot; N. L. adj. *philum* (from Gr. adj. *philos* -ê -on), friend,
 273 loving; N. L. adj. *thermophilum*, thermophilic).
 274

275 The description is based on three highly enriched monomethanogenic cultures. Cells are
 276 small, irregular, non-motile cocci, 0.4-0.5 µm. The cell wall is a thin monolayer covered with
 277 EPS. The cells lyze at salinity below 2 M NaCl. The F₄₂₀-dependent cell autofluorescence is
 278 absent. The core lipids are dominated by archaeol (C20-C20 DGE). The colony formation
 279 was not observed. Strictly anaerobic methanogens utilizing MeOH and trimethylamine as
 280 electron acceptor and H₂ or formate as electron donor. Heterotrophic, utilize yeast extract as
 281 C-source. The growth depends on external CoM and sterilized anaerobic sediments from salt
 282 lakes. Extremely halophilic, grow optimally at 4-5 M NaCl. The pH optimum for growth is 7-

283 7.5. Moderately thermophilic with an optimum at 50°C and the upper limit for growth at
 284 60°C. The G + C content of the genomic DNA in the type strain is 35.4 mol% (genome). The
 285 type strain, HMET1^T, was enriched from sediments of hypersaline lakes in Kulunda Steppe.
 286 The accession number of 16S rRNA sequence of the type strain in GenBank is KY449328.

287

288 **DESCRIPTION OF METHANONATRONARCHAEACEAE FAM. NOV.**

289 The description is the same as for the genus *Methanonatronarchaeum*.

290 Type genus: *Methanonatronarchaeum* gen. nov.

291

292 **DESCRIPTION OF ORDER METHANONATRONARCHAEALES ORD. NOV.**

293 The description is the same as for the genus *Methanonatronarchaeum*.

294 Type genus: *Methanonatronarchaeum* gen. nov.

295 **DESCRIPTION OF METHANONATRONARCHAEIA CLASSIS NOV.**

296 The class *Methanonatronarchaeia* is defined on the basis of comparative sequence analysis of
 297 the 16S rRNA obtained from 11 pure cultures of the genus *Methanonatronarchaeum*, 3 highly
 298 enriched cultures of 'Candidatus Methanohalarchaeum' and the cloned sequences from
 299 uncultured SA1 group found in various hypersaline habitats of terrestrial and marine origin.

300 Type order: *Methanonatronarchaeales* ord. nov.

301

302 **Fundings**

303 This work was supported by the Netherlands Applied Science Foundation (STW, project 12226),
 304 Gravitation (SIAM) grant 24002002 and by the Russian Foundation for Basic Research (RFBR 16-04-
 305 00035).

306

307 **Conflicts of interest**

308 The authors declared no conflicts of interest

309

310

311

312 **References**

313

- 314 1. **Oremland RS, King GM.** Methanogenesis in hypersaline environments. In: *Microbial*
 315 *mats. Physiological ecology of benthic microbial communities* 1989; pp 180-190. Cohen Y,
 316 Rosenberg E (eds). American Society for Microbiology: Washington, DC.
- 317 2. **McGenity TJ.** Methanogens and methanogenesis in hypersaline environments. In: Timmis
 318 KN (ed) *Handbook of Hydrocarbon and Lipid Microbiology* 2010; pp 665-679. Springer-
 319 Verlag: Berlin, Heidelberg.
- 320 3. **Paterek JR, Smith PH.** *Methanophilus mahii* gen. nov., sp. nov., a methylotrophic
 321 halophilic methanogen. *Int J Syst Bacteriol* 1988; 38: 122–123.
- 322 4. **Zhilina TN, Zavarzin GA.** *Methanohalobium evestigatus*, gen. nov. sp. nov., the extremely
 323 halophilic methanogenic archaeobacterium. *Dokl Akad Nauk USSR* 1987; 293: 464–468.

- 324 5. **Boone DR, Baker CC.** Genus VI. *Methanosalsum* gen. nov. In: *Bergey's Manual of*
325 *Systematic Bacteriology*, second edition 2001; vol 1: 287-289. Boone DR et al. (eds).
326 Springer-Verlag, New York.
- 327 6. **Mathrani JM, Boone DR, Mah RA, Fox GE, Lau PP.** *Methanohalophilus zhilinae* sp.
328 nov., an alkaliphilic, halophilic, methylotrophic methanogen. *Int J Syst Bacteriol* 1988; 38:
329 139–142.
- 330 7. **Kevbrin VV, Lysenko AM, Zhilina TN.** Physiology of alkaliphilic methanogen Z-7936, a
331 new strain of *Methanosalsus zhilinae* isolated from Lake Magadi. *Microbiology* (Moscow,
332 English Translation) 1997; 66: 261–266.
- 333 8. **Sorokin DY, Abbas BA, Sinninghe Damsté JS, Sukhacheva MV, van Loosdrecht MCM.**
334 *Methanocalculus alkaliphilus* sp. nov., and *Methanosalsum natronophilus* sp. nov., novel
335 haloalkaliphilic methanogens from hypersaline soda lakes. *Int J Syst Evol Microbiol* 2015;
336 65: 3739-3745.
- 337 9. **Menaia JAGF.** Osmotics of halophilic methanogenic archaeobacteria". *Scholar Archive*
338 *paper* 1992; 133 (<http://digitalcommons.ohsu.edu/etd>).
- 339 10. **Roberts MF, Lai MC, Gunsalus RP.** Biosynthetic pathways of the osmolytes Nε-acetyl-β-
340 lysine, β-glutamine, and betaine in *Methanohalophilus* strain FDF1 suggested by nuclear
341 magnetic resonance analyses. *J Bacteriol* 1992; 174: 6688–6693.
- 342 11. **Sorokin DY, Makarova K, Abbas B, Ferrer M, Golyshin PN.** Discovery of extremely
343 halophilic methyl-reducing methyl-reducing euryarchaea provides insights into the
344 evolutionary origin of methanogenesis. *Nature Microbiol* 2017; 2: article 17081.
- 345 12. **Fricke WF, Seedorf H, Henne A, Krüer M, Liesegang H.** The genome sequence of
346 *Methanosphaera stadtmanae* reveals why this human intestinal archaeon is restricted to
347 methanol and H₂ for methane formation and ATP synthesis. *J Bacteriol* 2006; 188: 642–
348 658.
- 349 13. **Sprenger WW, van Belzen MC, Rosenberg J, Hackstein JHP, Keltjens JT.**
350 *Methanomicrococcus blatticola* gen. nov., sp. nov., a methanol- and methylamine-reducing
351 methanogen from the hindgut of the cockroach *Periplaneta americana*. *Int J Syst Evol*
352 *Microbiol* 2000; 50: 1989-1999.
- 353 14. **Hedderich R, Whitman WB.** Physiology and biochemistry of the methane-producing
354 Archaea. In: *The Prokaryotes – Prokaryotic Physiology and Biochemistry* 2013; pp 635-
355 662. Rosenberg E et al. (eds). Springer-Verlag: Berlin, Heidelberg.
- 356 15. **Borrel G, Parisot N, Harris HMB, Peyretailade E, Gaci N et al.** Comparative genomics
357 highlights the unique biology of *Methanomassiliicoccales*, a *Thermoplasmatales*-related
358 seventh order of methanogenic archaea that encodes pyrrolysine. *BMC Genom* 2015; 15:
359 679.
- 360 16. **Paul K, Nonoh JO, Mikulski L, Brune A.** “Methanoplasmatales”, *Thermoplasmatales*-
361 related archaea in termite guts and other environments, are the seventh order of
362 methanogens. *Appl Environ Microbiol* 2012; 78: 8245-8253.

- 363 17. **Nobu MK, Narihiro T, Kuroda K, Mei R, Liu W-T.** Chasing the elusive *Euryarchaeota*
364 class WSA2: genomes reveal a uniquely fastidious methylreducing methanogen. *ISME J*
365 2016; 10: 2478-24-87.
- 366 18. **Evans PN, Parks DH, Chadwick GL, Robbins SJ, Orphan VJ.** Methane metabolism in
367 the archaeal phylum *Bathyarchaeota* revealed by genome-centric metagenomics. *Science*
368 2015; 350: 434-438.
- 369 19. **Vanwonterghem I, Evans PN, Parks DH, Jensen, PD, Woodcroft BJ.** Methylotrophic
370 methanogenesis discovered in the archaeal phylum *Verstraetearchaeota*. *Nat Microbiol*
371 2016; 1: article 16170.
- 372 20. **Weijers JWH, Panoto E, van Bleijswijk J, Schouten S, Balk M et al.** Constraints on the
373 biological source(s) of the orphan branched tetraether membrane lipids. *Geomicrobiol J*
374 2009; 26: 402-414.
- 375 21. **Sinninghe Damsté JS, Rijpstra WIC, Hopmans EC, Jung MY, Kim JG.** Intact polar and
376 core glycerol dibiphytanyl glycerol tetraether lipids of group I. 1a and I. 1b *Thaumarchaeota*
377 in soil. *Appl Environ Microbiol* 2012; 78: 6866-6874.
- 378 22. **Oremland RS, Marsh L, Desmarais DJ.** Methanogenesis in Big Soda Lake, Nevada: an
379 alkaline, moderately hypersaline desert lake. *Appl Environ Microbiol* 1982; 43, 462-468.
- 380 23. **Dawson KS, Freeman KH, Macalady JL.** Molecular characterization of core lipids from
381 halophilic archaea grown under different salinity conditions. *Org Chem* 2012; 48: 1-8.
- 382 24. **Guindon S, Dufayard JF, Lefort V, Anisimova M, Hordijk W et al.** New algorithms and
383 methods to estimate maximum-likelihood phylogenies: assessing the performance of
384 PhyML 3.0. *Syst Biol* 2010; 59: 307-21.
- 385 25. **Hordijk W, Gascuel O.** Improving the efficiency of SPR moves in phylogenetic tree search
386 methods based on maximum likelihood. *Bioinformatics* 2005; 21: 4338-47.
- 387 26. **Anisimova M, Gascuel O.** Approximate likelihood-ratio test for branches: A fast, accurate,
388 and powerful alternative. *Syst Biol* 2006; 55: 539-52.
- 389 27. **Quast C, Pruesse E, Yilmaz P, Gerken J, Schweer T et al.** The SILVA ribosomal RNA
390 gene database project: improved data processing and web-based tools. *Nucl Acids Res* 2013;
391 41: D590-D596.
- 392 28. **Eder W, Schmidt M, Koch M, Garbe-Schonberg D, Huber R.** Prokaryotic phylogenetic
393 diversity and corresponding geochemical data of the brine-seawater interface of the Shaban
394 Deep, Red Sea. *Environ microbiol* 2002; 4: 758-763.
- 395 29. **Ferrer M, Werner J, Chernikova TN, Bargiela R, Fernández L. et al.** Unveiling
396 microbial life in the new deep-sea hypersaline Lake *Thetis*. Part II: a metagenomic study.
397 *Environ Microbiol* 2012; 14: 268-281.
- 398 30. **Jiang H, Dong H, Yu B, Liu X, Li Y.** Microbial response to salinity change in Lake Chaka,
399 a hypersaline lake on Tibetan plateau. *Environ Microbiol* 2007; 9: 2603-2621.

400 **Table 1.** Extremely halophilic and moderately thermophilic mixotrophic methanogens isolated from hypersaline
401 lakes at 4 M total Na⁺ TMA - trimethylamine

Strain	Lake	Area	Brine parameters			Enrichment conditions		
			pH	Total salt g/l	Soluble carbonate alkalinity, M	Substrate	pH	T, °C
AMET1	Mix 5 soda lakes	Kulunda Steppe (Altai, Russia) 2013-2015	9.6-10.1	120-400	0.6-3.0	MeOH+formate	9.6	48
AMET3	Tanatar-1		10.1	350	2.8			48
AMET4	Picturesque Lake		9.8	250		48		
AMET5	Mix 6 soda lakes		9.6-10.2	50-380	0.5-3.4	TMA +formate		48
AMET6-2	Tanatar-1		10.25	380	3.4	MeOH+formate		60
AMET7	Soda crystallizer		9.6	350	3.8			55
AMET8	Mix 6 soda lakes		9.6-10.2	50-380				30
AMET9	Soda crystallizer		10.1	340	3.9			43
AMET10	Stamp Lake		9.1	325	0.2			54
AMET2	Mix from 8 lakes		Wadi al Natrun (Egypt, 2000)	9.1-9.9	200-360	0.1-0.9		MeOH+formate
AMET-SI	Searles Lake	California	9.8	350	0.2	MeOH+formate	9.2	48
HMET1 (mixed culture)	Mix from 4 salt lakes	Kulunda Steppe 2014	7.5-8.1	280-340	-	TMA+H ₂		48
HMET-EI (mixed culture)	Lake Elton	Southa Russia 2015	6.7	320	-	MeOH+formate	7.0	54
HMET-Eu (mixed culture)	Salt crystallizer	Crimea (Russia) 2015	7.2	220	-			55

402 **Figure legends**

403

404 **Fig. 1** Cell morphology of *Methanonatronarchaeum thermophilum* strain AMET1^T grown
405 with MeOH+formate at pH 9.5, 4 M total Na⁺ and 48°C. **(a)**, phase contrast microscopy; **(b**
406 **and c)**, electron microscopy of total cells and thin sections, respectively. **N**, nucleoid; **CM** -
407 cytoplasmic membrane; **CW** - cell wall.

408

409 **Fig. 2.** Cell morphology of '*Ca. Methanohalarchaeum thermophilum*' strain HMET1 grown
410 with TMA+H₂ at pH 7, 4 M NaCl and 50°C. **(a)**, phase contrast microscopy; **(b and c)**,
411 electron microscopy of total cells and thin section, respectively. **N**, nucleoid; **ICPM** -
412 intracytoplasmic membranes; **PHA** - possible polyhydroxyalkanoate storage granule; **CW** -
413 cell wall.

414

415 **Fig. 3.** Phylogeny of novel halo(alkali)philic methanogens from hypersaline lakes based on
416 the 16S rRNA gene sequence analysis. The bootstrap values above 70% are shown at the
417 nodes. Bar, 0.10 changes per position.

418

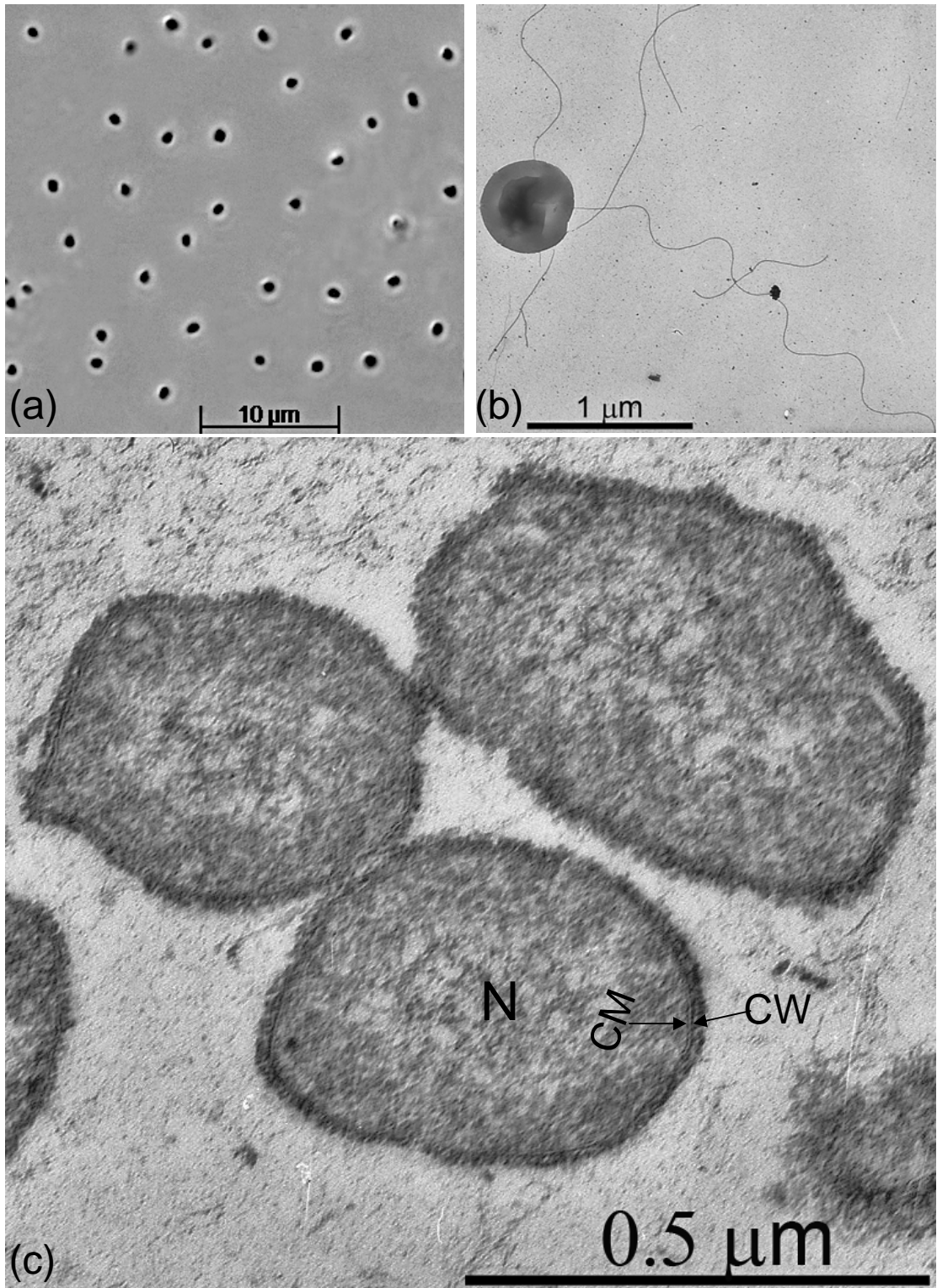


Fig.1

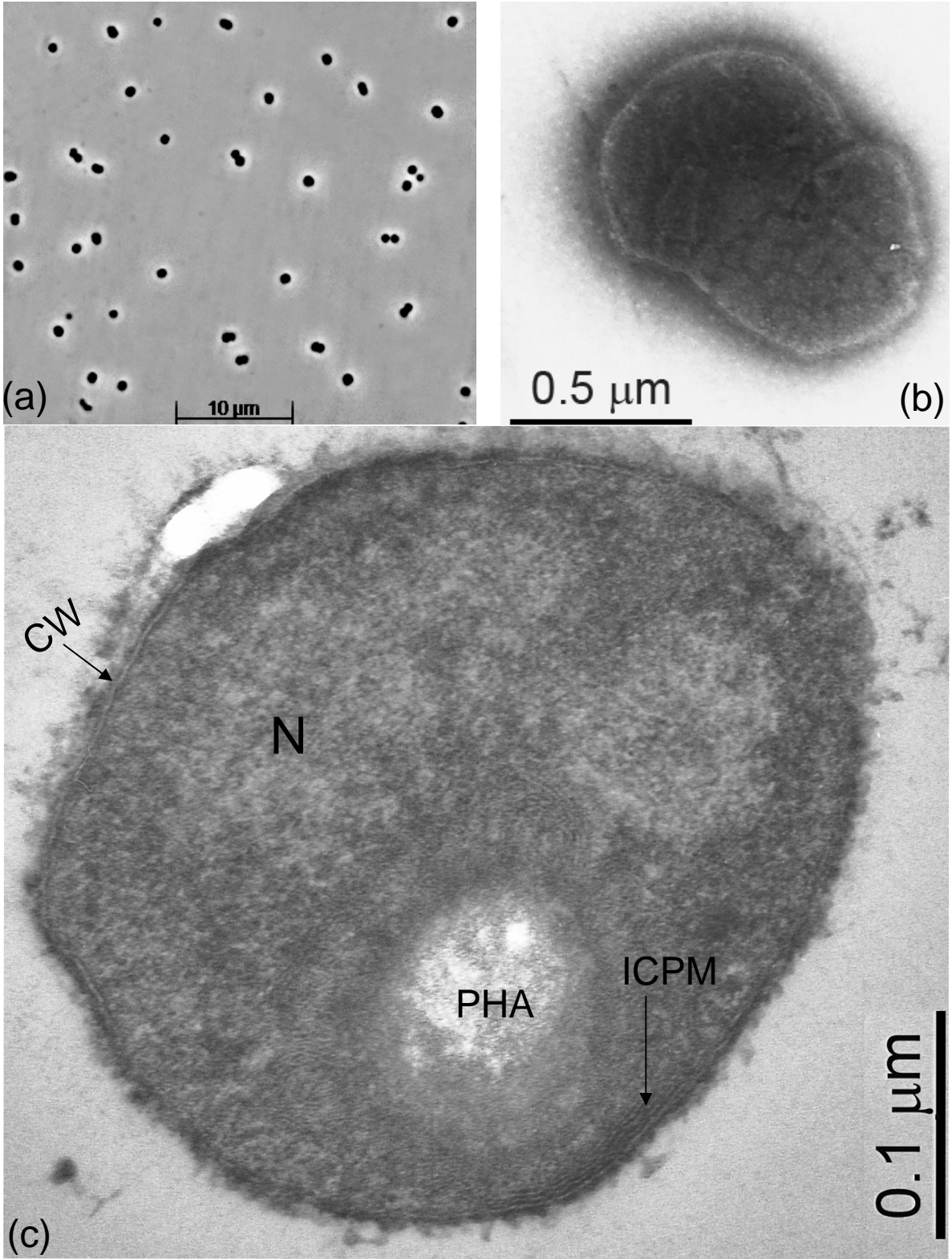
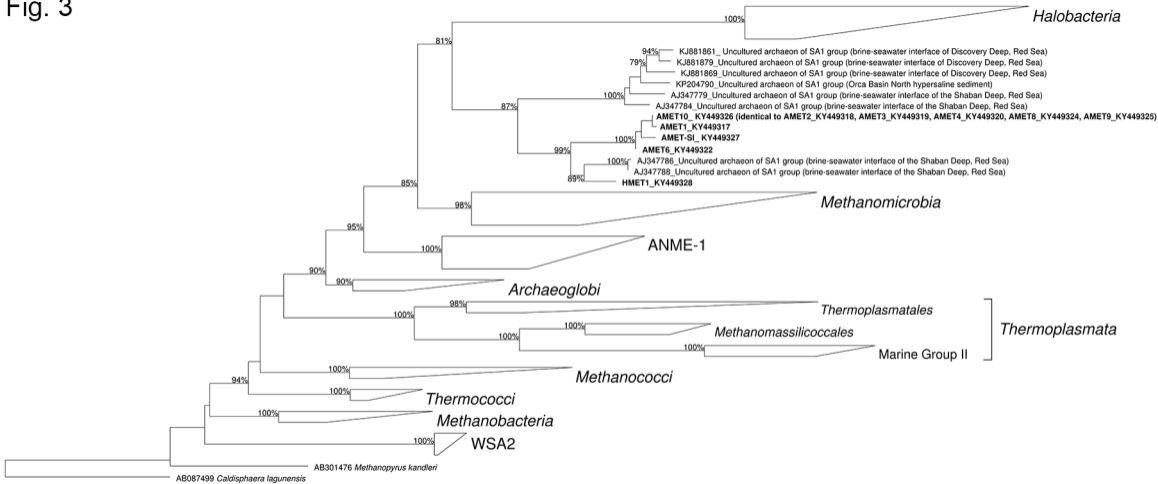


Fig.2

Fig. 3



***Methanonatronarchaeum thermophilum* gen. nov., sp. nov, and '*Candidatus Methanohalarchaeum thermophilum*', extremely halo(natrono)philic methyl-reducing methanogens from hypersaline lakes comprise a new euryarchaeal class *Methanonatronarchaeia* classis nov.**

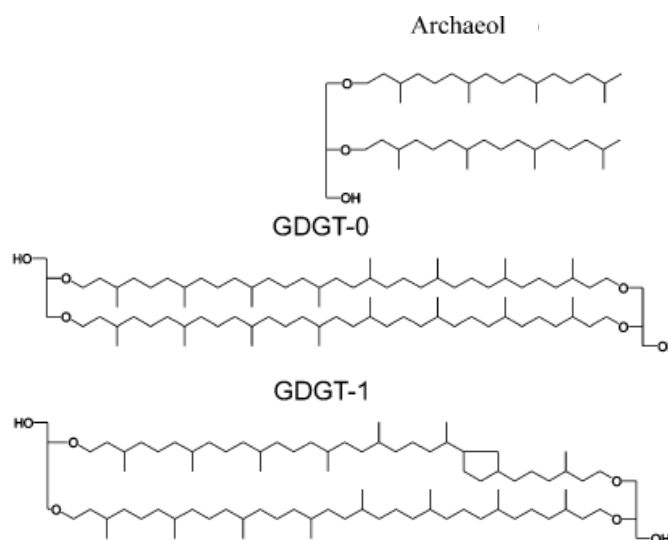
Dimitry Y. Sorokin^{a,b*}, Alexander Y. Merkel^a, Ben Abbas^b, Kira S. Makarova^c, W. Irene C. Rijpstra^d, M. Koenen^d, Jaap S. Sinninghe Damsté^{d,e}, Erwin A. Galinski^f, Eugene V. Koonin^c and Mark C.M. van Loosdrecht^b

Supplementary Table S1.

Membrane lipid composition of extremely halophilic methyl-reducing methanogens

A: core lipids

Strain	Di- and monophytanyl glycerol ethers (%)			Glycerol dibiphytanyl glycerol tetraethers	
	Archaeol	2-C20 MGE	1-C20 MGE	GDGT-0/archaeol	GDGT-1
AMET1	96.0	2.4	1.6	1:1	+
HMET1	93.4	-	6.6	3:1	-



B: Intact polar lipids composed of a core lipid with attached polar head group(s)

Strain	Archaeol							GDGT-0		
	PG	PGP	PGP-Me	PS	HPH	DH	Cardiolipin Ar-P-G-P- Ar*	PG PG	DH	
AMET1	++		+	+				++		
HMET1	+	+			+	+++	+		+++	

PG=Phosphatidylglycerol

PGP=Phosphatidylglycerolphosphate

PGP-Me=Phosphatidylglycerolphosphate methyl ester

PS=Phosphatidylserine

HPH=hexosephosphatidylhexose

DH=dihexose

Ar-P-G-P-Ar = archaeol-phosphatidyl-glycerol-phosphatidyl-archaeol

(*small part of the cardiolipin with extended archaeol (C25/C20) was also present)



RUSSIAN ACADEMY OF SCIENCES
WINOGRADSKY INSTITUTE OF MICROBIOLOGY
RESEARCH CENTRE OF BIOTECHNOLOGY

117312 Russia, Moscow, Prospekt 60-let Oktyabrya, 7/2
Tel. (095) 135-21-39; Fax (095) 135-65-30; e-mail: inmi@inmi.ru

Moscow, Russia

12 September 2014

Confirmation of the availability of a strain for the purpose of valid publication of a new name according to the Bacteriological Code

The following information is confidential and serves only to allow the Microbiology journal to confirm that a strain has been deposited and will be available from the UNIQEM (Unique and Extremophilic Microorganisms Collection of Winogradsky Institute of Microbiology RAS).

Methanonatronaerchaeium thermophilum" strain **AMET1(T)** has been deposited in the UNIQEM under the number **U982**.

The strains are available in the open section of the UNIQEM and restrictions have not been placed on access to information concerning the presence of the strain in the UNIQEM. It will be included in published and online catalogues after publication of this number by the authors.

The strain has been checked for viability and is stored using 20% glycerol stock culture in liquid nitrogen.

Prof. Dr. Valery F. Galchenko
Director of UNIQEM