



## Strain identifier

**BacDive ID:** 7219      **DOI:** 10.13145/bacdive7219.20190402.4  
**Type strain:** yes      **Designation:** H2s  
**Culture col. no.:** DSM 21852, VKM B-2545

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## Name and taxonomic classification

<a href="#">Ref.: 15991</a>	<b>Domain</b>	Bacteria
<a href="#">Ref.: 15991</a>	<b>Phylum</b>	Proteobacteria
<a href="#">Ref.: 15991</a>	<b>Class</b>	Alphaproteobacteria
<a href="#">Ref.: 15991</a>	<b>Order</b>	Rhizobiales
<a href="#">Ref.: 15991</a>	<b>Family</b>	Methylocystaceae
<a href="#">Ref.: 15991</a>	<b>Genus</b>	Methylocystis
<a href="#">Ref.: 15991</a>	<b>Species</b>	Methylocystis bryophila
<a href="#">Ref.: 15991</a>	<b>Full Scientific Name</b>	Methylocystis bryophila Belova et al. 2013
<a href="#">Ref.: 15991</a>	<b>Designation:</b>	H2s
<a href="#">Ref.: 15991</a>	<b>Type strain:</b>	yes

### **Prokaryotic Nomenclature Up-to-date (PNU)**

<a href="#">Ref.: 20215</a>	<b>Domain</b>	Bacteria
<a href="#">Ref.: 20215</a>	<b>Phylum</b>	Proteobacteria
<a href="#">Ref.: 20215</a>	<b>Class</b>	Alphaproteobacteria
<a href="#">Ref.: 20215</a>	Literature reference	Int. J. Syst. Evol. Microbiol. 56:1
<a href="#">Ref.: 20215</a>	<b>Family</b>	Methylocystaceae
<a href="#">Ref.: 20215</a>	<b>Genus</b>	Methylocystis
<a href="#">Ref.: 20215</a>	Taxonomical status	gen. nov. (VP)

Ref.: 20215	Literature reference	Int. J. Syst. Bacteriol. 43:735*
Ref.: 20215	<b>Species</b>	Methylocystis bryophila
Ref.: 20215	Taxonomical status	sp. nov. (VP)
Ref.: 20215	Literature reference	Int. J. Syst. Evol. Microbiol. 63:1096*
Ref.: 20215	<b>Full Scientific Name</b>	Methylocystis bryophila Belova et al. 2013

### Morphology and physiology

Ref.: 30743	<b>Gram stain</b>	negative
Ref.: 30743	<b>Cell length</b>	2.6 µm
Ref.: 30743	<b>Cell width</b>	1.15 µm
Ref.: 30743	<b>Cell shape</b>	coccus-shaped
Ref.: 30743	<b>Motility</b>	no

	<b>Metabolite utilization</b>	<b>Chebi ID</b>	<b>Metabolite</b>	<b>Utilization activity</b>	<b>Kind of utilization tested</b>
Ref.: 30743		16449	Alanine	+	carbon source
Ref.: 30743		22653	Asparagine	+	carbon source
Ref.: 30743		18237	Glutamic acid	+	carbon source
Ref.: 30743		26271	Proline	+	carbon source
Ref.: 30743		17822	Serine	+	carbon source

Ref.: 15991	<b>Oxygen tolerance</b>	aerobe
Ref.: 30743	<b>Oxygen tolerance</b>	aerobe
Ref.: 30743	<b>Ability of spore formation</b>	no

### Culture and growth conditions

Ref.: 15991	<b>Culture medium</b>	METHYLOCYSTIS MEDIUM (DSMZ Medium 1409), 22°C, pH 5.8 - 6.2; aerobic with 10-20% methane in the headspace
Ref.: 15991	<b>Culture medium growth</b>	yes
Ref.: 15991	<b>Culture medium link</b>	<a href="https://www.dsmz.de/microorganisms/medium/pdf/DSMZ_Medium1409.pdf">https://www.dsmz.de/microorganisms/medium/pdf/DSMZ_Medium1409.pdf</a>

	<b>Temperatures</b>	<b>Kind of temperature</b>	<b>Temperature</b>
Ref.: 15991		growth	20-25 °C
Ref.: 30743		growth	08-37 °C
Ref.: 30743		optimum	27.5 °C



Ref.: 30743      **Temperature range**      mesophilic

Ref.: 30743 Ref.: 30743	<b>pH</b>	<b>Kind of pH</b>	<b>pH</b>
		growth	4.2-7.6
		optimum	6.25

**Isolation, sampling and environmental information**

Ref.: 15991      **Sample type/isolated from**      Sphagnum peat, bank of a bog lake

Ref.: 15991      **Host species**      Sphagnum

Ref.: 15991      **Geographic location (country and/or sea, region)**      north eastern Germany, lake Teufelssee

Ref.: 15991      **Country**      Germany

Ref.: 15991      **Continent**      Europe

<b>Isolation sources categories</b>	<b>Cat1</b>	<b>Cat2</b>	<b>Cat3</b>
	#Host	#Plants	#Peat moss

**Application and interaction**

Ref.: 15991      **Biosafety level**      1 Risk group (German classification)

**Molecular biology**

Ref.: 15991      **GC-content**      62 mol% thermal denaturation, midpoint method (Tm)

Ref.: 30743      **GC-content**      62 mol%

	<b>Sequence database</b>	<b>Sequence accession description</b>	<b>Sequence accession number</b>	<b>Sequence length(bp)</b>	<b>Associated NCBI tax ID</b>
Ref.: 15991	EMBL Direct submission		FN422004		
Ref.: 15991	EMBL Direct submission		FN422005		
Ref.: 15991	EMBL Direct submission	Methylocystis bryophila partial pmoA2 gene for particulate methane monooxygenase active-site polypeptide, type strain H2sT	HE798546	531	655015



<a href="#">Ref.: 15991</a>	EMBL Direct submission	Methylocystis bryophila partial nifH gene for nitrogenase iron protein, type strain H2sT	HE798545	449	655015
<a href="#">Ref.: 15991</a>	EMBL Direct submission		FN422003		

## Strain availability

[Ref.: 15991](#)      **Culture collection no.**      DSM 21852, VKM B-2545

[Ref.: 15991](#)      **Strain history**      <- S. N. Dedysh, Winogradski Inst. Microbiol., RAS, Moscow, Russia; H2s <- S. E. Belova and S. N. Dedysh

### **Associated Passport(s) in StrainInfo**

[Ref.: 20218](#)      885193 - <http://www.straininfo.net/strains/885193>

[Ref.: 20218](#)      885194 - <http://www.straininfo.net/strains/885194>

## References

[Ref.: 15991](#)      Leibniz Institut DSMZ-Deutsche Sammlung von Mikroorganismen und Zellkulturen GmbH; Curators of the DSMZ; DSM 21852

[Ref.: 20215](#)      D.Gleim, M.Kracht, N.Weiss et. al.: Prokaryotic Nomenclature Up-to-date - compilation of all names of Bacteria and Archaea, validly published according to the Bacteriological Code since 1. Jan. 1980, and validly published nomenclatural changes since.

[Ref.: 20218](#)      Verslyppe, B., De Smet, W., De Baets, B., De Vos, P., Dawyndt P. StrainInfo introduces electronic passports for microorganisms.. Syst Appl Microbiol. 37: 42-50 2014 (10.1016/j.syapm.2013.11.002, 24321274)

[Ref.: 30743](#)      Barberan A, Caceres Velazquez H, Jones S, Fierer N. Hiding in Plain Sight: Mining Bacterial Species Records for Phenotypic Trait Information. mSphere 2: None-None 2017 (10.1128/mSphere.00237-17, None) - **originally annotated from #27074**

[Ref.: 27074](#)      IJSEM 1096 2013 (10.1099/ijms.0.043505-0)

**\* These References are textmined**

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