

**Taxonomic Outline of the Bacteria and Archaea, Release 7.7 March 6, 2007.
Part 6– Phylum “*Proteobacteria*”: The *Deltaproteobacteria* and
*Epsilonproteobacteria***

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Class *Deltaproteobacteria*^{VP} Kuever et al 2006. N4Lid DOI: 10.1601/nm.3456

Order *Desulfurellales*^{VP} Kuever et al 2006. N4Lid DOI: 10.1601/nm.3457

Family *Desulfurellaceae*^{VP} Kuever et al 2006. N4Lid DOI: 10.1601/nm.3458

Genus *Desulfurella*^{VP} Bonch-Osmolovskaya et al. 1993. N4Lid DOI: 10.1601/nm.3459

Desulfurella acetivorans^{VP} Bonch-Osmolovskaya et al. 1993. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 51451. High-quality 16S rRNA sequence S000008874 (RDP), X72768 (Genbank). N4Lid DOI: 10.1601/nm.3460

Desulfurella kamchatkensis^{VP} Miroshnichenko 1998. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 700655. High-quality 16S rRNA sequence S000000228 (RDP), Y16941 (Genbank). N4Lid DOI: 10.1601/nm.3461

Desulfurella multipotens^{VP} Miroshnichenko et al. 1996. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 8415. High-quality 16S rRNA sequence S000004736 (RDP), Y16943 (Genbank). N4Lid DOI: 10.1601/nm.3462

Desulfurella propionica^{VP} Miroshnichenko 1998. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 700656. High-quality 16S rRNA sequence S000007307 (RDP), Y16942 (Genbank). N4Lid DOI: 10.1601/nm.3463

Genus *Hippea*^{VP} Miroshnichenko et al. 1999. N4Lid DOI: 10.1601/nm.3464

Hippea maritima^{VP} Miroshnichenko et al. 1999. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 700847. High-quality 16S rRNA sequence S000016943 (RDP), Y18292 (Genbank). N4Lid DOI: 10.1601/nm.3465

Order *Desulfovibrionales*^{VP} Kuever et al 2006. N4Lid DOI: 10.1601/nm.3466

Family *Desulfovibrionaceae*^{VP} Kuever et al 2006. N4Lid DOI: 10.1601/nm.3467

Genus *Desulfovibrio*^{AL} Kluyver and van Niel 1936. ≈ *Desulfovibrio* (later heterotypic synonym), links through 10.1601/nm.3468. N4Lid DOI: 10.1601/nm.3468

Desulfovibrio desulfuricans *subsp. desulfuricans*^{AL} (Beijerinck 1895) Kluyver and van Niel 1936. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 29577. High-quality 16S rRNA sequence S000006150 (RDP), AF192153 (Genbank). GOLD ID: Gc00315. GCAT ID: 000295_GCAT. Entrez genome id: 329. Sequenced strain: G20 is from a non-type strain. Genome sequencing is completed. Number of genomes of this species sequenced 1 (GOLD). N4Lid DOI: 10.1601/nm.3469

Desulfovibrio desulfuricans *subsp. aestuarii*^{AL} Postgate and Campbell 1966. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 29578. N4Lid DOI: 10.1601/nm.3470

Desulfovibrio acrylicus^{VP} van der Maarel et al. 1997. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 10141. High-quality 16S rRNA sequence S000437480 (RDP), U32578 (Genbank). N4Lid DOI: 10.1601/nm.3471

Desulfovibrio aespoensis^{VP} Motamedi and Pedersen 1998. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 700646. High-quality 16S rRNA sequence S000112712 (RDP), X95230 (Genbank). N4Lid DOI: 10.1601/nm.3472

Desulfovibrio africanus^{AL} Campbell et al. 1966. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 19996. High-quality 16S rRNA sequence S000003900 (RDP), X99236 (Genbank). N4Lid DOI: 10.1601/nm.3473

Desulfovibrio alaskensis^{VP} Feio et al. 2004. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 16109. High-quality 16S rRNA sequence S000560277 (RDP), Y11984 (Genbank). N4Lid DOI: 10.1601/nm.8570

- Desulfovibrio alcoholivorans*^{VP} corrig. Qatibi et al. 1995. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 49738. High-quality 16S rRNA sequence S000428080 (RDP), AF050101 (Genbank). N4Lid DOI: 10.1601/nm.3474
- Desulfovibrio alkalitolerans*^{VP} Abildgaard et al. 2006. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 16529. High-quality 16S rRNA sequence S000421932 (RDP), AY649785 (Genbank). N4Lid DOI: 10.1601/nm.10047
- Desulfovibrio aminophilus*^{VP} Baena et al. 1999. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 12254. High-quality 16S rRNA sequence S000428372 (RDP), AF067964 (Genbank). N4Lid DOI: 10.1601/nm.3475
- Desulfovibrio bastinii*^{VP} Magot et al. 2004. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-903. High-quality 16S rRNA sequence S000461354 (RDP), AY359868 (Genbank). N4Lid DOI: 10.1601/nm.8571
- Desulfovibrio burkinensis*^{VP} Ouattara et al. 1999. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 700846. High-quality 16S rRNA sequence S000416204 (RDP), AF053752 (Genbank). N4Lid DOI: 10.1601/nm.3478
- Desulfovibrio carbinolicus*^{VP} Nanninga and Gottschal 1995. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-1241. N4Lid DOI: 10.1601/nm.3479
- Desulfovibrio cuneatus*^{VP} Sass et al. 1998. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 11391. N4Lid DOI: 10.1601/nm.3480
- Desulfovibrio dechloracetivorans*^{VP} Sun et al. 2001. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 700912. High-quality 16S rRNA sequence S000389681 (RDP), AF230530 (Genbank). N4Lid DOI: 10.1601/nm.3481
- Desulfovibrio ferrireducens*^{VP} Vandieken et al. 2006. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 16995. High-quality 16S rRNA sequence S000588198 (RDP), DQ148945 (Genbank). N4Lid 10.1601/nm.9984
- Desulfovibrio frigidus*^{VP} Vandieken et al. 2006. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 17176. High-quality 16S rRNA sequence S000588196 (RDP), DQ148943 (Genbank). N4Lid 10.1601/nm.9985
- Desulfovibrio fructosivorans*^{VP} corrig. Ollivier et al. 1990. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 49200. N4Lid DOI: 10.1601/nm.3482
- Desulfovibrio furfuralis*^{VP} Folkerts et al. 1989. Source of type material recommended for DOE sponsored genome sequencing by the JGI: F1. N4Lid DOI: 10.1601/nm.3483
- Desulfovibrio gabonensis*^{VP} Tardy-Jacquenod et al. 1996. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 700201. High-quality 16S rRNA sequence S000437467 (RDP), U31080 (Genbank). N4Lid DOI: 10.1601/nm.3484
- Desulfovibrio giganteus*^{VP} Esnault et al. 1988. Synonym links through N4Lid: 10.1601/nm.3485. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 4123. N4Lid DOI: 10.1601/nm.3485
- † *Desulfobacter giganteus*^{VP} Esnault et al. 1988. Orthographic error. *Desulfobacter giganteus*. Links through N4Lid 10.1601/8. > Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 4123. N4Lid DOI: 10.1601/nm.3543
- Desulfovibrio gigas*^{AL} Le Gall 1963. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 19364. High-quality 16S rRNA sequence S000436230 (RDP), M34400 (Genbank). N4Lid DOI: 10.1601/nm.3486
- Desulfovibrio gracilis*^{VP} Magot et al. 2004. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-904. N4Lid 10.1601/nm.8956
- Desulfovibrio halophilus*^{VP} Caumette et al. 1991. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 51179. High-quality 16S rRNA sequence S000011580 (RDP), X99237 (Genbank). N4Lid DOI: 10.1601/nm.3487
- Desulfovibrio hydrothermalis*^{VP} Alazard et al. 2003. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 14728. High-quality 16S rRNA sequence S000394495 (RDP), AF458778 (Genbank). N4Lid DOI: 10.1601/nm.3488

Desulfovibrio indonesiensis^{VP} corrig. Feio et al. 2000. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 15121. High-quality 16S rRNA sequence S000539470 (RDP), AJ621884 (Genbank). N4Lid DOI: 10.1601/nm.3489

Desulfovibrio inopinatus^{VP} Reichenbecher and Schink 1999. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 10711. High-quality 16S rRNA sequence S000010738 (RDP), AF177276 (Genbank). N4Lid DOI: 10.1601/nm.3490

Desulfovibrio intestinalis^{VP} Fröhlich et al. 1999. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 11275. High-quality 16S rRNA sequence S000016406 (RDP), Y12254 (Genbank). N4Lid DOI: 10.1601/nm.3491

Desulfovibrio litoralis^{VP} Sass et al. 1998. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 11393. N4Lid DOI: 10.1601/nm.3492

Desulfovibrio longreachensis^{VP} corrig. Redburn and Patel 1995. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ACM 3958. High-quality 16S rRNA sequence S000130468 (RDP), Z24450 (Genbank). N4Lid DOI: 10.1601/nm.3493

Desulfovibrio longus^{VP} Magot et al. 1992. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 51456. High-quality 16S rRNA sequence S000461353 (RDP), AY359867 (Genbank). N4Lid DOI: 10.1601/nm.3494

Desulfovibrio magneticus^{VP} Sakaguchi et al. 2002. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 700980. High-quality 16S rRNA sequence S000413803 (RDP), D43944 (Genbank). GOLD ID: Gi00729. GCAT ID: 001001_GCAT. Entrez genome id: 15681. Sequenced strain: ATCC 700980 is type material. Genome sequencing is incomplete. Number of genomes of this species sequenced 1 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3495

Desulfovibrio mexicanus^{VP} Hernandez-Eugenio et al. 2001. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 13116. High-quality 16S rRNA sequence S000010910 (RDP), AF227984 (Genbank). N4Lid DOI: 10.1601/nm.3496

Desulfovibrio oxamicus^{VP} (Postgate and Campbell 1966) López-Cortés et al. 2006. <== *Desulfovibrio vulgaris subsp. oxamicus* (basonym). Synonym links through N4Lid: 10.1601/ex.3509. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 33405. High-quality 16S rRNA sequence S000584779 (RDP), DQ122124 (Genbank). N4Lid 10.1601/nm.9781

† *Desulfovibrio vulgaris subsp. oxamicus*^{AL} Postgate and Campbell 1966. ==> *Desulfovibrio oxamicus* (new combination). Synonym links through N4Lid: 10.1601/ex.3509. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 33405. High-quality 16S rRNA sequence S000584779 (RDP), DQ122124 (Genbank). N4Lid DOI: 10.1601/nm.3509

Desulfovibrio oxyclinae^{VP} Krekeler et al. 2000. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 11498. N4Lid DOI: 10.1601/nm.3497

Desulfovibrio piger^{VP} (Moore et al. 1976) Loubinoux et al. 2002. <== *Desulfomonas pigra* (basonym). Synonym links through N4Lid: 10.1601/ex.3527. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 29098. High-quality 16S rRNA sequence S000001126 (RDP), AF192152 (Genbank). GOLD ID: Gi01734. GCAT ID: 000680_GCAT. Sequenced strain: ATCC 29098 is type material. Genome sequencing is incomplete. Number of genomes of this species sequenced 1 (GOLD). N4Lid 10.1601/nm.9219

† *Desulfomonas pigra*^{AL} Moore et al. 1976. ==> *Desulfovibrio piger* (new combination). Synonym links through N4Lid: 10.1601/ex.3527. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 29098. High-quality 16S rRNA sequence S000001126 (RDP), AF192152 (Genbank). GOLD ID: Gi01734. GCAT ID: 000680_GCAT. Sequenced strain: ATCC 29098 is type material. Genome sequencing is incomplete. Number of genomes of this species sequenced 1 (GOLD) 0 (NCBI). N4Lid DOI: 10.1601/nm.3527

"*Desulfovibrio pigra*" † <==Synonym links through N4Lid: 10.1601/ex.3527. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 29098. High-quality 16S rRNA sequence S000001126 (RDP), AF192152 (Genbank). GOLD ID: Gi01734. GCAT ID: 000680_GCAT. Sequenced strain: ATCC 29098 is type material. Genome sequencing is incomplete. Number of genomes of this species sequenced 1 (GOLD) 0 (NCBI). N4Lid DOI: 10.1601/nm.3498

Desulfovibrio profundus^{VP} Bale et al. 1997. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 11384. High-quality 16S rRNA sequence S000438770 (RDP), U90726 (Genbank). N4Lid DOI: 10.1601/nm.3499

Desulfovibrio putealis^{VP} Basso et al. 2005. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-905. High-quality 16S rRNA sequence S000475288 (RDP), AY574979 (Genbank). N4Lid DOI: 10.1601/nm.8573

Desulfovibrio salexigens^{AL} Postgate and Campbell 1966. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 14822. High-quality 16S rRNA sequence S000436231 (RDP), M34401 (Genbank). N4Lid DOI: 10.1601/nm.3500

Desulfovibrio sapovorans^{VP} Widdel 1981. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 33892. High-quality 16S rRNA sequence S000436232 (RDP), M34402 (Genbank). N4Lid DOI: 10.1601/nm.3501

"*Desulfovibrio sapovorans*" † *Desulfovibrio sapovorans* (invalid name), links through 10.1601/nm.3501. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 33892. High-quality 16S rRNA sequence S000436232 (RDP), M34402 (Genbank). N4Lid DOI: 10.1601/nm.3564

Desulfovibrio senezii^{VP} Tsu et al. 1999. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 8436. High-quality 16S rRNA sequence S000428079 (RDP), AF050100 (Genbank). N4Lid DOI: 10.1601/nm.3502

Desulfovibrio simplex^{VP} Zellner et al. 1990. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 4141. N4Lid DOI: 10.1601/nm.3503

Desulfovibrio sulfodismutans^{VP} Bak and Pfennig 1988. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 43913. High-quality 16S rRNA sequence S000012355 (RDP), Y17764 (Genbank). N4Lid DOI: 10.1601/nm.3504

Desulfovibrio termidis^{VP} Trinkerl et al. 1991. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 49858. High-quality 16S rRNA sequence S000004739 (RDP), X87409 (Genbank). N4Lid DOI: 10.1601/nm.3505

Desulfovibrio vietnamensis^{VP} Dang et al. 2002. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 10520. High-quality 16S rRNA sequence S000083867 (RDP), X93994 (Genbank). N4Lid DOI: 10.1601/nm.3507

Desulfovibrio vulgaris subsp. vulgaris^{AL} Postgate and Campbell 1966. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 29579. High-quality 16S rRNA sequence S000436229 (RDP), M34399 (Genbank). GOLD ID: Gc00184. GCAT ID: 000426_GCAT. Entrez genome id: 51. Sequenced strain: ATCC 29579 is type material. Genome sequencing is completed. Number of genomes of this species sequenced 2 (GOLD) 2 (NCBI). N4Lid DOI: 10.1601/nm.3508

Desulfovibrio zosteriae^{VP} Nielsen et al. 1999. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 11974. High-quality 16S rRNA sequence S000020196 (RDP), Y18049 (Genbank). N4Lid DOI: 10.1601/nm.3510

Genus † *Desulfomonas*^{AL} Moore et al. 1976. ≈ *Desulfovibrio* (earlier heterotypic synonym), links through 10.1601/nm.3468. N4Lid DOI: 10.1601/nm.3526

Genus *Bilophila*^{VP} Baron et al. 1990. N4Lid DOI: 10.1601/nm.3511

Bilophila wadsworthia^{VP} Baron et al. 1990. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 49260. N4Lid DOI: 10.1601/nm.3512

Genus *Lawsonia*^{VP} McOrist et al. 1995. N4Lid DOI: 10.1601/nm.3513

Lawsonia intracellularis^{VP} McOrist et al. 1995. Source of type material recommended for DOE sponsored genome sequencing by the JGI: NCTC 12656. High-quality 16S rRNA sequence S000437459 (RDP), U30147 (Genbank). GOLD ID: Gc00382. GCAT ID: 000228_GCAT. Entrez genome id: 183. Sequenced strain: NCTC 12657 is from a non-type strain. Genome sequencing is completed. Number of genomes of this species sequenced 1 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3514

Family *Desulfomicrobiaceae*^{VP} Kuever et al 2006. N4Lid DOI: 10.1601/nm.3515

Genus *Desulfomicrobium*^{VP} Rozanova et al. 1994. N4Lid DOI: 10.1601/nm.3516

Desulfomicrobium baculatum^{VP} corrig. (Rozanova and Nazina 1984) Rozanova et al. 1994. <== *Desulfovibrio baculatus* (basonym). Synonym links through N4Lid: 10.1601/ex.3477. Source of

type material recommended for DOE sponsored genome sequencing by the JGI: DSM 4028. High-quality 16S rRNA sequence S000427865 (RDP), AF030438 (Genbank). N4Lid DOI: 10.1601/nm.3517

† *Desulfovibrio baculatus*^{VP} Rozanova and Nazina 1984. ==> *Desulfomicrobium baculatum* (new combination). Synonym links through N4Lid: 10.1601/ex.3477. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 4028. High-quality 16S rRNA sequence S000427865 (RDP), AF030438 (Genbank). N4Lid DOI: 10.1601/nm.3477

Desulfomicrobium apsheronum^{VP} Rozanova et al. 1994. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 5918. High-quality 16S rRNA sequence S000438109 (RDP), U64865 (Genbank). N4Lid DOI: 10.1601/nm.3518

Desulfomicrobium escambiense^{VP} corrig. Sharak Genthner et al. 1996. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 51164. High-quality 16S rRNA sequence S000436976 (RDP), U02469 (Genbank). N4Lid DOI: 10.1601/nm.3519

Desulfomicrobium macestii^{VP} (Gogotova and Vainshtein 1989) Hippe et al. 2003. <==> *Desulfobacterium macestii* (basonym). Synonym links through N4Lid: 10.1601/ex.3556. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 4194. High-quality 16S rRNA sequence S000137299 (RDP), AJ237604 (Genbank). N4Lid DOI: 10.1601/nm.3520

† *Desulfobacterium macestii*^{VP} Gogotova and Vainstein 1989. ==> *Desulfomicrobium macestii* (new combination). Synonym links through N4Lid: 10.1601/ex.3556. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 4194. High-quality 16S rRNA sequence S000137299 (RDP), AJ237604 (Genbank). N4Lid DOI: 10.1601/nm.3556

Desulfomicrobium norvegicum^{VP} Genthner et al. 1997. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 1741. High-quality 16S rRNA sequence S000138146 (RDP), AJ277897 (Genbank). N4Lid DOI: 10.1601/nm.3521

Desulfomicrobium orale^{VP} Langendijk et al. 2001. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 12838. High-quality 16S rRNA sequence S000008660 (RDP), AJ251623 (Genbank). N4Lid DOI: 10.1601/nm.3522

Family *Desulfhalobiaceae*^{VP} Kuever et al. 2006. N4Lid DOI: 10.1601/nm.3523

Genus *Desulfhalobium*^{VP} Ollivier et al. 1991. N4Lid DOI: 10.1601/nm.3524

Desulfhalobium retbaense^{VP} Ollivier et al. 1991. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 49708. High-quality 16S rRNA sequence S000437689 (RDP), U48244 (Genbank). N4Lid DOI: 10.1601/nm.3525

Desulfhalobium utahense^{VP} Jakobsen et al. 2006. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 17720. High-quality 16S rRNA sequence S000556967 (RDP), DQ067421 (Genbank). N4Lid DOI: 10.1601/nm.10281

Genus *Desulfonatronovibrio*^{VP} Zhilina et al. 1997. N4Lid DOI: 10.1601/nm.3528

Desulfonatronovibrio hydrogenovorans^{VP} Zhilina et al. 1997. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 9292. High-quality 16S rRNA sequence S000002343 (RDP), X99234 (Genbank). N4Lid DOI: 10.1601/nm.3529

Genus *Desulfonauticus*^{VP} Audiffren et al. 2003. N4Lid DOI: 10.1601/nm.3530

Desulfonauticus submarinus^{VP} Audiffren et al. 2003. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 15269. High-quality 16S rRNA sequence S000431412 (RDP), AF524933 (Genbank). N4Lid DOI: 10.1601/nm.3531

Genus *Desulfothermus*^{VP} Kuever et al. 2006. N4Lid DOI: 10.1601/nm.3532

Desulfothermus naphthae^{VP} Kuever et al. 2006. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 13418. High-quality 16S rRNA sequence S000010081 (RDP), X80922 (Genbank). N4Lid DOI: 10.1601/nm.3533

Family "*Desulfonatronaceae*" N4Lid DOI: 10.1601/nm.3534

Genus *Desulfonatronum*^{VP} Pikuta et al. 1998. N4Lid DOI: 10.1601/nm.3535

Desulfonatronum lacustre^{VP} Pikuta et al. 1998. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 10312. High-quality 16S rRNA sequence S000003106 (RDP), Y14594 (Genbank). N4Lid DOI: 10.1601/nm.3536

Desulfonatronum cooperativum^{VP} Zhilina et al. 2005. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 16749. High-quality 16S rRNA sequence S000549637 (RDP), AY725424 (Genbank). N4Lid 10.1601/nm.9421

Desulfonatronum thiodismutans^{VP} Pikuta et al. 2003. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-395. High-quality 16S rRNA sequence S000539197 (RDP), AF373920 (Genbank). N4Lid DOI: 10.1601/nm.3537

Order Desulfobacterales^{VP} Kuever et al 2006. N4Lid DOI: 10.1601/nm.3538

Family Desulfobacteraceae^{VP} Kuever et al 2006. N4Lid DOI: 10.1601/nm.3539

Genus Desulfobacter^{VP} Widdel 1981. N4Lid DOI: 10.1601/nm.3540

Desulfobacter postgatei^{VP} Widdel 1981. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 33911. High-quality 16S rRNA sequence S000393859 (RDP), AF418180 (Genbank). N4Lid DOI: 10.1601/nm.3541

Desulfobacter curvatus^{VP} Widdel 1988. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 43919. High-quality 16S rRNA sequence S000436243 (RDP), M34413 (Genbank). N4Lid DOI: 10.1601/nm.3542

Desulfobacter halotolerans^{VP} Brandt and Ingvorsen 1998. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 11383. High-quality 16S rRNA sequence S000003902 (RDP), Y14745 (Genbank). N4Lid DOI: 10.1601/nm.3544

Desulfobacter hydrogenophilus^{VP} Widdel 1988. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 43915. High-quality 16S rRNA sequence S000436242 (RDP), M34412 (Genbank). N4Lid DOI: 10.1601/nm.3545

Desulfobacter latus^{VP} Widdel 1988. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 43918. High-quality 16S rRNA sequence S000436244 (RDP), M34414 (Genbank). N4Lid DOI: 10.1601/nm.3546

Desulfobacter vibrioformis^{VP} Lien and Beeder 1997. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 8776. High-quality 16S rRNA sequence S000437115 (RDP), U12254 (Genbank). N4Lid DOI: 10.1601/nm.3547

Genus Desulfatibacillum^{VP} Cravo-Laureau et al. 2004. N4Lid DOI: 10.1601/nm.3548

Desulfatibacillum aliphaticivorans^{VP} Cravo-Laureau et al. 2004. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-743. High-quality 16S rRNA sequence S000400333 (RDP), AY184360 (Genbank). N4Lid DOI: 10.1601/nm.3549

Desulfatibacillum alkenivorans^{VP} Cravo-Laureau et al. 2004. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-924. High-quality 16S rRNA sequence S000420138 (RDP), AY493562 (Genbank). N4Lid DOI: 10.1601/nm.8569

Genus Desulfobacterium^{VP} Bak and Widdel 1988. N4Lid DOI: 10.1601/nm.3550

Desulfobacterium indolicum^{VP} Bak and Widdel 1988. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 43938. High-quality 16S rRNA sequence S000088673 (RDP), AJ237607 (Genbank). N4Lid DOI: 10.1601/nm.3551

Desulfobacterium anilini^{VP} Schnell et al. 1990. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 49792. High-quality 16S rRNA sequence S000137583 (RDP), AJ237601 (Genbank). N4Lid DOI: 10.1601/nm.3552

Desulfobacterium autotrophicum^{VP} Brysch et al. 1988. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 43914. High-quality 16S rRNA sequence S000436239 (RDP), M34409 (Genbank). GOLD ID: Gi00194. GCAT ID: 000998_GCAT. Entrez genome id: 109. Sequenced strain: HRM2 is type material. Genome sequencing is incomplete. Number of genomes of this species sequenced 1 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3553

Desulfobacterium catecholicum^{VP} Szewzyk and Pfennig 1988. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 43955. High-quality 16S rRNA sequence S000089909 (RDP), AJ237602 (Genbank). N4Lid DOI: 10.1601/nm.3554

Desulfobacterium niacini^{VP} Kuever et al. 2006. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 2650. N4Lid DOI: 10.1601/nm.3557

Desulfobacterium vacuolatum^{VP} Kuever et al. 2006. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 3385. High-quality 16S rRNA sequence S000436238 (RDP), M34408 (Genbank). N4Lid DOI: 10.1601/nm.3559

- Genus *Desulfobacula*** ^{VP} Rabus et al. 2000. N4Lid DOI: 10.1601/nm.3560
- Desulfobacula toluolica* ^{VP} Rabus et al. 2000. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 7467. High-quality 16S rRNA sequence S000112434 (RDP), X70953 (Genbank). N4Lid DOI: 10.1601/nm.3561
- Desulfobacula phenolica* ^{VP} (Bak and Widdel 1988) Kuever et al. 2001. <== *Desulfobacterium phenolicum* (basonym). Synonym links through N4Lid: 10.1601/ex.3558. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 43956. High-quality 16S rRNA sequence S000137584 (RDP), AJ237606 (Genbank). N4Lid DOI: 10.1601/nm.3562
- † *Desulfobacterium phenolicum* ^{VP} Bak and Widdel 1988. ==> *Desulfobacula phenolica* (new combination). Synonym links through N4Lid: 10.1601/ex.3558. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 43956. High-quality 16S rRNA sequence S000137584 (RDP), AJ237606 (Genbank). N4Lid DOI: 10.1601/nm.3558
- Genus "*Desulfobotulus*"** † Invalid name. DOI: 10.1601/nm.3563
- Genus *Desulfocella*** ^{VP} Brandt et al. 1999. N4Lid DOI: 10.1601/nm.3565
- Desulfocella halophila* ^{VP} Brandt et al. 1999. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 700426. High-quality 16S rRNA sequence S000427693 (RDP), AF022936 (Genbank). N4Lid DOI: 10.1601/nm.3566
- Genus *Desulfococcus*** ^{VP} Widdel 1981. N4Lid DOI: 10.1601/nm.3567
- Desulfococcus multivorans* ^{VP} Widdel 1981. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 33890. High-quality 16S rRNA sequence S000436235 (RDP), M34405 (Genbank). N4Lid DOI: 10.1601/nm.3568
- Desulfococcus biacutus* ^{VP} Platen et al. 1991. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 5651. High-quality 16S rRNA sequence S000018513 (RDP), AJ277887 (Genbank). N4Lid DOI: 10.1601/nm.3569
- Genus *Desulfofaba*** ^{VP} Knoblauch et al. 1999. ≈ *Desulfofaba* (later heterotypic synonym), links through 10.1601./nm.3570. N4Lid DOI: 10.1601/nm.3570
- Desulfofaba gelida* ^{VP} Knoblauch et al. 1999. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 12344. High-quality 16S rRNA sequence S000428876 (RDP), AF099063 (Genbank). N4Lid DOI: 10.1601/nm.3571
- Desulfofaba fastidiosa* ^{VP} Abildgaard et al. 2004. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-815. High-quality 16S rRNA sequence S000441050 (RDP), AY268891 (Genbank). N4Lid DOI: 10.1601/nm.3572
- Desulfofaba hansenii* ^{VP} (Finster et al. 2001) Abildgaard et al. 2004. <== *Desulfomusa hansenii* (basonym). Synonym links through N4Lid: 10.1601/ex.3578. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 700811. High-quality 16S rRNA sequence S000391502 (RDP), AF321820 (Genbank). N4Lid DOI: 10.1601/nm.3573
- † *Desulfomusa hansenii* ^{VP} Finster et al. 2001. ==> *Desulfofaba hansenii* (new combination). Synonym links through N4Lid: 10.1601/ex.3578. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 700811. High-quality 16S rRNA sequence S000391502 (RDP), AF321820 (Genbank). N4Lid DOI: 10.1601/nm.3578
- Genus † *Desulfomusa*** ^{VP} Finster et al. 2001. ≈ *Desulfofaba* (earlier heterotypic synonym), links through 10.1601./nm.3570. N4Lid DOI: 10.1601/nm.3577
- Genus *Desulfofrigus*** ^{VP} Knoblauch et al. 1999. N4Lid DOI: 10.1601/nm.3574
- Desulfofrigus oceanense* ^{VP} Knoblauch et al. 1999. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 12341. High-quality 16S rRNA sequence S000428877 (RDP), AF099064 (Genbank). N4Lid DOI: 10.1601/nm.3575
- Desulfofrigus fragile* ^{VP} Knoblauch et al. 1999. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 12345. High-quality 16S rRNA sequence S000428878 (RDP), AF099065 (Genbank). N4Lid DOI: 10.1601/nm.3576
- Genus *Desulfonema*** ^{VP} Widdel 1981. N4Lid DOI: 10.1601/nm.3579
- Desulfonema limicola* ^{VP} Widdel 1981. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 33961. High-quality 16S rRNA sequence S000437653 (RDP), U45990 (Genbank). N4Lid DOI: 10.1601/nm.3580

Desulfonema ishimotonii^{VP} corrig. Fukui et al. 2000. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 9680. High-quality 16S rRNA sequence S000437655 (RDP), U45992 (Genbank). N4Lid DOI: 10.1601/nm.3581

Desulfonema magnum^{VP} Widdel 1981. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 35288. High-quality 16S rRNA sequence S000437652 (RDP), U45989 (Genbank). N4Lid DOI: 10.1601/nm.3582

Genus *Desulforegula*^{VP} Rees and Patel 2001. N4Lid DOI: 10.1601/nm.3583

Desulforegula conservatrix^{VP} Rees and Patel 2001. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-134. High-quality 16S rRNA sequence S000389920 (RDP), AF243334 (Genbank). N4Lid DOI: 10.1601/nm.3584

Genus *Desulfosarcina*^{VP} Widdel 1981. N4Lid DOI: 10.1601/nm.3585

Desulfosarcina variabilis^{VP} Widdel 1981. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 33932. High-quality 16S rRNA sequence S000436147 (RDP), M26632 (Genbank). N4Lid DOI: 10.1601/nm.3586

Desulfosarcina cetonica^{VP} corrig. (Galushko and Rozanova 1994) Kuever et al. 2006. <== *Desulfobacterium cetonicum* (basonym). Synonym links through N4Lid: 10.1601/ex.3555. Source of type material recommended for DOE sponsored genome sequencing by the JGI: JCM 12296. High-quality 16S rRNA sequence S000088251 (RDP), AJ237603 (Genbank). N4Lid 10.1601/nm.9792

† *Desulfobacterium cetonicum*^{VP} Galushko and Rozanova 1994. ==> *Desulfosarcina cetonica* (new combination). Synonym links through N4Lid: 10.1601/ex.3555. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 7267. High-quality 16S rRNA sequence S000088251 (RDP), AJ237603 (Genbank). N4Lid DOI: 10.1601/nm.3555

"*Desulfosarcina cetonium*" Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 7267. High-quality 16S rRNA sequence S000088251 (RDP), AJ237603 (Genbank). N4Lid DOI: 10.1601/nm.3587

Desulfosarcina ovata^{VP} Kuever et al. 2006. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 13228. N4Lid DOI: 10.1601/nm.3588

Genus *Desulfospira*^{VP} Finster et al. 1997. N4Lid DOI: 10.1601/nm.3589

Desulfospira joergensenii^{VP} Finster et al. 1997. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 700409. High-quality 16S rRNA sequence S000015339 (RDP), X99637 (Genbank). N4Lid DOI: 10.1601/nm.3590

Genus *Desulfotignum*^{VP} Kuever et al. 2001. N4Lid DOI: 10.1601/nm.3591

Desulfotignum balticum^{VP} Kuever et al. 2001. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-19. High-quality 16S rRNA sequence S000389726 (RDP), AF233370 (Genbank). N4Lid DOI: 10.1601/nm.3592

Desulfotignum phosphitoxidans^{VP} Schink et al. 2002. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 13687. High-quality 16S rRNA sequence S000393881 (RDP), AF420288 (Genbank). N4Lid DOI: 10.1601/nm.3593

Family *Desulfobulbaceae*^{VP} Kuever et al 2006. N4Lid DOI: 10.1601/nm.3594

Genus *Desulfobulbus*^{VP} Widdel 1981. N4Lid DOI: 10.1601/nm.3595

Desulfobulbus propionicus^{VP} Widdel 1981. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 33891. High-quality 16S rRNA sequence S000436240 (RDP), M34410 (Genbank). N4Lid DOI: 10.1601/nm.3596

Desulfobulbus elongatus^{VP} Samain et al. 1985. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 43118. High-quality 16S rRNA sequence S000126615 (RDP), X95180 (Genbank). N4Lid DOI: 10.1601/nm.3597

Desulfobulbus mediterraneus^{VP} Sass et al. 2002. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 13871. High-quality 16S rRNA sequence S000392134 (RDP), AF354663 (Genbank). N4Lid DOI: 10.1601/nm.3598

Desulfobulbus rhabdiformis^{VP} Lien et al. 1998. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 700652. High-quality 16S rRNA sequence S000437114 (RDP), U12253 (Genbank). N4Lid DOI: 10.1601/nm.3599

Genus *Desulfocapsa*^{VP} Janssen et al. 1997. N4Lid DOI: 10.1601/nm.3600

Desulfocapsa thiozymogenes^{VP} Janssen et al. 1997. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 7269. High-quality 16S rRNA sequence S000128257 (RDP), X95181 (Genbank). N4Lid DOI: 10.1601/nm.3601

Desulfocapsa sulfexigens^{VP} corrig. Finster et al. 2000. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 700424. High-quality 16S rRNA sequence S000130725 (RDP), Y13672 (Genbank). N4Lid DOI: 10.1601/nm.3602

Genus *Desulfofustis*^{VP} Friedrich et al. 1996. N4Lid DOI: 10.1601/nm.3603

Desulfofustis glycolicus^{VP} Friedrich et al. 1996. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 700454. High-quality 16S rRNA sequence S000381437 (RDP), X99707 (Genbank). N4Lid DOI: 10.1601/nm.3604

Genus *Desulforhopalus*^{VP} Isaksen and Teske 1999. N4Lid DOI: 10.1601/nm.3605

Desulforhopalus vacuolatus^{VP} Isaksen and Teske 1999. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 700480. High-quality 16S rRNA sequence S000435986 (RDP), L42613 (Genbank). N4Lid DOI: 10.1601/nm.3606

Desulforhopalus singaporensis^{VP} Lie et al. 2000. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 12130. High-quality 16S rRNA sequence S000429262 (RDP), AF118453 (Genbank). N4Lid DOI: 10.1601/nm.3607

Genus *Desulfotalea*^{VP} Knoblauch et al. 1999. N4Lid DOI: 10.1601/nm.3608

Desulfotalea psychrophila^{VP} Knoblauch et al. 1999. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 12343. High-quality 16S rRNA sequence S000428875 (RDP), AF099062 (Genbank). GOLD ID: Gc00207. GCAT ID: 000023_GCAT. Entrez genome id: 12751. Sequenced strain: DSM 12343 is type material. Genome sequencing is completed. Number of genomes of this species sequenced 1 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3609

Desulfotalea arctica^{VP} Knoblauch et al. 1999. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 12342. High-quality 16S rRNA sequence S000428874 (RDP), AF099061 (Genbank). N4Lid DOI: 10.1601/nm.3610

Family *Nitrospinaceae*^{VP} Garrity et al 2006. N4Lid DOI: 10.1601/nm.3611

Genus *Nitrospina*^{AL} Watson and Waterbury 1971. N4Lid DOI: 10.1601/nm.3612

Nitrospina gracilis^{AL} Watson and Waterbury 1971. Source of type material recommended for DOE sponsored genome sequencing by the JGI: No viable type strain.. High-quality 16S rRNA sequence S000414629 (RDP), L35504 (Genbank). GOLD ID: Gi00860. GCAT ID: 001273_GCAT. Entrez genome id: 13445. Sequenced strain: Nb 211 is from a non-type strain. Genome sequencing is incomplete. Number of genomes of this species sequenced 1 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3613

Order *Desulfarculales*^{VP} N4Lid DOI: 10.1601/nm.3614

Family *Desulfarculaceae*^{VP} Kuever et al 2006. N4Lid DOI: 10.1601/nm.3615

Genus *Desulfarculus*^{VP} Kuever et al. 2006. N4Lid DOI: 10.1601/nm.3616

Desulfarculus baarsii^{VP} (Widdel 1981) Kuever et al. 2006. <== *Desulfovibrio baarsii* (basonym). Synonym links through N4Lid: 10.1601/ex.3476. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 33931. High-quality 16S rRNA sequence S000436233 (RDP), M34403 (Genbank). N4Lid DOI: 10.1601/nm.3617

† *Desulfovibrio baarsii*^{VP} Widdel 1981. ==> *Desulfarculus baarsii* (new combination). Synonym links through N4Lid: 10.1601/ex.3476. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 33931. High-quality 16S rRNA sequence S000436233 (RDP), M34403 (Genbank). N4Lid DOI: 10.1601/nm.3476

Order *Desulfuromonadales*^{VP} Kuever et al. 2006. N4Lid DOI: 10.1601/nm.3618

Family *Desulfuromonadaceae*^{VP} Kuever et al. 2006. N4Lid DOI: 10.1601/nm.3619

Genus *Desulfuromonas*^{AL} Pfennig and Biebl 1977. N4Lid DOI: 10.1601/nm.3620

Desulfuromonas acetoxidans^{AL} Pfennig and Biebl 1977. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 684. High-quality 16S rRNA sequence S000436149 (RDP), M26634 (Genbank). GOLD ID: Gi00197. GCAT ID: 001002_GCAT. Entrez genome id: 12551. Sequenced strain: DSM 684 is type material. Genome sequencing is

incomplete. Number of genomes of this species sequenced 1 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3621

Desulfuromonas acetexigens^{VP} Finster et al. 1997. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 1397. N4Lid DOI: 10.1601/nm.3622

Desulfuromonas chloroethenica^{VP} Krumholz 1997. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 700295. High-quality 16S rRNA sequence S000381082 (RDP), U49748 (Genbank). N4Lid DOI: 10.1601/nm.3623

Desulfuromonas palmitatis^{VP} Coates et al. 2000. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 51701. High-quality 16S rRNA sequence S000437417 (RDP), U28172 (Genbank). N4Lid DOI: 10.1601/nm.3624

Desulfuromonas svalbardensis^{VP} Vandieken et al. 2006. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 16958. High-quality 16S rRNA sequence S000457290 (RDP), AY835388 (Genbank). N4Lid DOI: 10.1601/nm.10052

Desulfuromonas thiophila^{VP} Finster et al. 1997. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 700367. High-quality 16S rRNA sequence S000007761 (RDP), Y11560 (Genbank). N4Lid DOI: 10.1601/nm.3625

Genus *Desulfuromusa*^{VP} Liesack and Finster 1994. N4Lid DOI: 10.1601/nm.3626

Desulfuromusa kysingii^{VP} Liesack and Finster 1994. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 7343. High-quality 16S rRNA sequence S000015699 (RDP), X79414 (Genbank). N4Lid DOI: 10.1601/nm.3627

Desulfuromusa bakii^{VP} Liesack and Finster 1994. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 7345. High-quality 16S rRNA sequence S000016405 (RDP), X79412 (Genbank). N4Lid DOI: 10.1601/nm.3628

Desulfuromusa ferrireducens^{VP} Vandieken et al. 2006. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 16956. High-quality 16S rRNA sequence S000457294 (RDP), AY835392 (Genbank). N4Lid DOI: 10.1601/nm.10053

Desulfuromusa succinoxidans^{VP} Liesack and Finster 1994. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 8270. High-quality 16S rRNA sequence S000006970 (RDP), X79415 (Genbank). N4Lid DOI: 10.1601/nm.3629

Genus *Malonomonas*^{VP} Dehning and Schink 1990. N4Lid DOI: 10.1601/nm.3630

Malonomonas rubra^{VP} Dehning and Schink 1990. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 5091. High-quality 16S rRNA sequence S000010833 (RDP), Y17712 (Genbank). N4Lid DOI: 10.1601/nm.3631

Genus *Pelobacter*^{VP} Schink and Pfennig 1983. N4Lid DOI: 10.1601/nm.3632

Pelobacter acidigallici^{VP} Schink and Pfennig 1983. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 49970. High-quality 16S rRNA sequence S000003147 (RDP), X77216 (Genbank). N4Lid DOI: 10.1601/nm.3633

Pelobacter acetylenicus^{VP} Schink 1986. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 3246. High-quality 16S rRNA sequence S000009676 (RDP), X70955 (Genbank). N4Lid DOI: 10.1601/nm.3634

Pelobacter carbinolicus^{VP} Schink 1984. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 2380. High-quality 16S rRNA sequence S000008920 (RDP), X79413 (Genbank). GOLD ID: Gc00306. GCAT ID: 000304_GCAT. Entrez genome id: 13337. Sequenced strain: DSM 2380 is type material. Genome sequencing is completed. Number of genomes of this species sequenced 1 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3635

Pelobacter massiliensis^{VP} Schnell et al. 1991. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 49973. N4Lid DOI: 10.1601/nm.3636

Pelobacter propionicus^{VP} Schink 1984. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 2379. High-quality 16S rRNA sequence S000005137 (RDP), X70954 (Genbank). GOLD ID: Gc00467. GCAT ID: 000142_GCAT. Entrez genome id: 13384. Sequenced strain: DSM 2379 is type material. Genome sequencing is completed. Number of genomes of this species sequenced 1 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3637

Pelobacter venetianus^{VP} Schink and Stieb 1984. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 35615. High-quality 16S rRNA sequence S000437600 (RDP), U41562 (Genbank). N4Lid DOI: 10.1601/nm.3638

Family Geobacteraceae^{VP} Holmes et al 2004. N4Lid DOI: 10.1601/nm.3639

Genus Geobacter^{VP} Lovley et al. 1995. GOLD ID: Gi00944. GCAT ID: 001102_GCAT. Entrez genome id: 16263. Sequenced strain: FRC-32 is from a non-type strain. Genome sequencing is incomplete. Number of genomes of this species sequenced 3 (GOLD) 3 (NCBI). N4Lid DOI: 10.1601/nm.3640

Geobacter metallireducens^{VP} Lovley et al. 1995. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 53774. High-quality 16S rRNA sequence S000414324 (RDP), L07834 (Genbank). GOLD ID: Gc00314. GCAT ID: 000296_GCAT. Entrez genome id: 177. Sequenced strain: GS-15 is from a non-type strain. Genome sequencing is completed. Number of genomes of this species sequenced 1 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3641

Geobacter bemidjensis^{VP} Nevin et al. 2005. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-1014. High-quality 16S rRNA sequence S000571800 (RDP), AY187307 (Genbank). GOLD ID: Gi01650. GCAT ID: 001103_GCAT. Entrez genome id: 17707. Sequenced strain: ATCC BAA-1014 is type material. Genome sequencing is incomplete. Number of genomes of this species sequenced 1 (GOLD) 1 (NCBI). N4Lid 10.1601/nm.9537

Geobacter brementis^{VP} Straub and Buchholz-Cleven 2001. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-607. High-quality 16S rRNA sequence S000438915 (RDP), U96917 (Genbank). N4Lid DOI: 10.1601/nm.3642

Geobacter chapellei^{VP} Coates et al. 2001. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 51744. High-quality 16S rRNA sequence S000437599 (RDP), U41561 (Genbank). N4Lid DOI: 10.1601/nm.3643

Geobacter grbciae^{VP} Coates et al. 2001. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-45. High-quality 16S rRNA sequence S000391801 (RDP), AF335182 (Genbank). N4Lid DOI: 10.1601/nm.3644

Geobacter hydrogenophilus^{VP} Coates et al. 2001. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 51590. High-quality 16S rRNA sequence S000437418 (RDP), U28173 (Genbank). N4Lid DOI: 10.1601/nm.3645

Geobacter pelophilus^{VP} Straub and Buchholz-Cleven 2001. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-603. High-quality 16S rRNA sequence S000438916 (RDP), U96918 (Genbank). N4Lid DOI: 10.1601/nm.3646

Geobacter psychrophilus^{VP} Nevin et al. 2005. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-1013. High-quality 16S rRNA sequence S000548805 (RDP), AY653548 (Genbank). N4Lid 10.1601/nm.9538

Geobacter sulfurreducens^{VP} Caccavo et al. 1995. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 51573. High-quality 16S rRNA sequence S000437158 (RDP), U13928 (Genbank). GOLD ID: Gc00166. GCAT ID: 000444_GCAT. Entrez genome id: 192. Sequenced strain: ATCC 51573 is type material. Genome sequencing is completed. Number of genomes of this species sequenced 1 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3647

Genus Geopsychrobacter^{VP} Holmes et al. 2005. N4Lid 10.1601/nm.9283

Geopsychrobacter electrodiphilus^{VP} Holmes et al. 2005. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-880. High-quality 16S rRNA sequence S000440962 (RDP), AY187303 (Genbank). N4Lid 10.1601/nm.9282

Genus Geothermobacter^{VP} Kashefi et al. 2005. N4Lid 10.1601/nm.9285

Geothermobacter ehrlichii^{VP} Kashefi et al. 2005. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-635 and DSM 15274. High-quality 16S rRNA sequence S000398534 (RDP), AY155599 (Genbank). N4Lid 10.1601/nm.9284

Genus Trichlorobacter^{VP} De Wever et al. 2001. N4Lid DOI: 10.1601/nm.3648

Trichlorobacter thiogenes^{VP} De Wever et al. 2001. *Geobacter thiogenes* (new combination). Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC

BAA-34. High-quality 16S rRNA sequence S000389424 (RDP), AF223382 (Genbank). N4Lid DOI: 10.1601/nm.3649

Order Syntrophobacterales^{VP} Kuever et al 2006. N4Lid DOI: 10.1601/nm.3650

Family Syntrophobacteraceae^{VP} Kuever et al 2006. N4Lid DOI: 10.1601/nm.3651

Genus Syntrophobacter^{VP} Boone and Bryant 1984. N4Lid DOI: 10.1601/nm.3652

Syntrophobacter wolinii^{VP} Boone and Bryant 1984. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 2805. High-quality 16S rRNA sequence S000004801 (RDP), X70905 (Genbank). N4Lid DOI: 10.1601/nm.3653

Syntrophobacter fumaroxidans^{VP} Harmsen et al. 1998. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 10017. High-quality 16S rRNA sequence S000022640 (RDP), X82874 (Genbank). GOLD ID: Gc00453. GCAT ID: 000156_GCAT. Entrez genome id: 13013. Sequenced strain: MPOB is type material. Genome sequencing is completed. Number of genomes of this species sequenced 1 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3654

Syntrophobacter pfennigii^{VP} Wallrabenstein et al. 1996. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 10092. High-quality 16S rRNA sequence S000017105 (RDP), X82875 (Genbank). N4Lid DOI: 10.1601/nm.3655

Syntrophobacter sulfatireducens^{VP} Chen et al. 2005. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 16706. High-quality 16S rRNA sequence S000548711 (RDP), AY651787 (Genbank). N4Lid DOI: 10.1601/nm.9452

Genus Desulfacinum^{VP} Rees et al. 1995. N4Lid DOI: 10.1601/nm.3656

Desulfacinum infernum^{VP} Rees et al. 1995. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 9756. High-quality 16S rRNA sequence S000414554 (RDP), L27426 (Genbank). N4Lid DOI: 10.1601/nm.3657

Desulfacinum hydrothermale^{VP} Sievert and Kuever 2000. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 13146. High-quality 16S rRNA sequence S000388514 (RDP), AF170417 (Genbank). N4Lid DOI: 10.1601/nm.3658

Genus Desulforhabdus^{VP} Oude Elferink et al. 1997. N4Lid DOI: 10.1601/nm.3659

Desulforhabdus amnigena^{VP} corrig. Oude Elferink et al. 1997. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 51979. High-quality 16S rRNA sequence S000010018 (RDP), X83274 (Genbank). N4Lid DOI: 10.1601/nm.3660

Genus Desulfovira^{VP} Tanaka et al. 2000. N4Lid DOI: 10.1601/nm.3661

Desulfovira adipica^{VP} Tanaka et al. 2000. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 12016. N4Lid DOI: 10.1601/nm.3662

Genus Thermodesulforhabdus^{VP} Beeder et al. 1996. N4Lid DOI: 10.1601/nm.3663

Thermodesulforhabdus norvegica^{VP} corrig. Beeder et al. 1996. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 51998. High-quality 16S rRNA sequence S000437365 (RDP), U25627 (Genbank). N4Lid DOI: 10.1601/nm.3664

Family Syntrophaceae^{VP} Kuever et al 2006. N4Lid DOI: 10.1601/nm.3665

Genus Syntrophus^{VP} Mountfort et al. 1984. N4Lid DOI: 10.1601/nm.3666

Syntrophus buswellii^{VP} Mountfort et al. 1984. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 2612A. High-quality 16S rRNA sequence S000015757 (RDP), X85131 (Genbank). N4Lid DOI: 10.1601/nm.3667

Syntrophus aciditrophicus^{VP} Jackson et al. 2001. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 700169. High-quality 16S rRNA sequence S000438631 (RDP), U86447 (Genbank). GOLD ID: Gc00341. GCAT ID: 000269_GCAT. Entrez genome id: 16258. Sequenced strain: SB is from a non-type strain. Genome sequencing is completed. Number of genomes of this species sequenced 2 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3668

Syntrophus gentianae^{VP} Wallrabenstein et al. 1996. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 8423. High-quality 16S rRNA sequence S000004383 (RDP), X85132 (Genbank). N4Lid DOI: 10.1601/nm.3669

Genus Desulfobacca^{VP} Oude Elferink et al. 1999. N4Lid DOI: 10.1601/nm.3670

Desulfobacca acetoxidans^{VP} Oude Elferink et al. 1999. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 700848. High-quality 16S rRNA sequence S000386891 (RDP), AF002671 (Genbank). N4Lid DOI: 10.1601/nm.3671

Genus *Desulfomonile*^{VP} DeWeerd et al. 1991. N4Lid DOI: 10.1601/nm.3672

Desulfomonile tiedjei^{VP} DeWeerd et al. 1991. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 49306. High-quality 16S rRNA sequence S000436150 (RDP), M26635 (Genbank). N4Lid DOI: 10.1601/nm.3673

Desulfomonile limimaris^{VP} Sun et al. 2001. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 700979. High-quality 16S rRNA sequence S000389682 (RDP), AF230531 (Genbank). N4Lid DOI: 10.1601/nm.3674

Genus *Smithella*^{VP} Liu et al. 1999. N4Lid DOI: 10.1601/nm.3675

Smithella propionica^{VP} Liu et al. 1999. Source of type material recommended for DOE sponsored genome sequencing by the JGI: OCM 661. High-quality 16S rRNA sequence S000387293 (RDP), AF126282 (Genbank). N4Lid DOI: 10.1601/nm.3676

Order *Bdellovibrionales*^{VP} Garrity et al 2006. N4Lid DOI: 10.1601/nm.3677

Family *Bdellovibrionaceae*^{VP} Garrity et al 2006. N4Lid DOI: 10.1601/nm.3678

Genus *Bdellovibrio*^{AL} Stolp and Starr 1963. N4Lid DOI: 10.1601/nm.3679

Bdellovibrio bacteriovorus^{AL} Stolp and Starr 1963. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 15356. High-quality 16S rRNA sequence S000130039 (RDP), AJ292759 (Genbank). GOLD ID: Gc00168. GCAT ID: 000442_GCAT. Entrez genome id: 9637. Sequenced strain: ATCC 15356 is type material. Genome sequencing is completed. Number of genomes of this species sequenced 2 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3680

Genus *Micavibrio*^{VP} Lambina et al. 1989. N4Lid DOI: 10.1601/nm.3686

Micavibrio admirandus^{VP} Lambina et al. 1989. Source of type material recommended for DOE sponsored genome sequencing by the JGI: VKM B-1619. N4Lid DOI: 10.1601/nm.3687

Genus *Vampirovibrio*^{VP} Gromov and Mamkayeva 1980. N4Lid DOI: 10.1601/nm.3688

Vampirovibrio chlorellavorus^{VP} (ex Gromov and Mamkayeva 1972) Gromov and Mamkayeva 1980. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 29753. N4Lid DOI: 10.1601/nm.3689

Family *Bacteriovoraceae*^{VP} Davidov and Jurkevitch 2004. N4Lid 10.1601/nm.8976

Genus *Bacteriovorax*^{VP} Baer et al. 2000. N4Lid DOI: 10.1601/nm.3683

Bacteriovorax stolpii^{VP} (Seidler et al. 1972) Baer et al. 2000. <== *Bdellovibrio stolpii* (**basonym**). Synonym links through N4Lid: 10.1601/ex.3682. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 27052. High-quality 16S rRNA sequence S000436219 (RDP), M34125 (Genbank). N4Lid DOI: 10.1601/nm.3684

† *Bdellovibrio stolpii*^{AL} Seidler et al. 1972. ==> *Bacteriovorax stolpii* (**new combination**). Synonym links through N4Lid: 10.1601/ex.3682. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 27052. High-quality 16S rRNA sequence S000436219 (RDP), M34125 (Genbank). N4Lid DOI: 10.1601/nm.3682

Bacteriovorax litoralis^{VP} Baer et al. 2004. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-684. High-quality 16S rRNA sequence S000365693 (RDP), AF084859 (Genbank). N4Lid DOI: 10.1601/nm.8507

Bacteriovorax marinus^{VP} Baer et al. 2004. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-682. High-quality 16S rRNA sequence S000365691 (RDP), AF084854 (Genbank). GOLD ID: Gi00066. GCAT ID: 000763_GCAT. Entrez genome id: 12622. Sequenced strain: SJ is type material. Genome sequencing is incomplete. Number of genomes of this species sequenced 1 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.8508

Genus *Peredibacter*^{VP} Davidov and Jurkevitch 2004. N4Lid 10.1601/nm.8769

Peredibacter starrii^{VP} (Seidler et al. 1972) Davidov and Jurkevitch 2004. <== *Bdellovibrio starrii* (**basonym**). Synonym links through N4Lid: 10.1601/ex.3681. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 15145. High-quality 16S rRNA sequence S000428666 (RDP), AF084852 (Genbank). N4Lid 10.1601/nm.8770

† *Bacteriovorax starrii*^{VP} (Seidler et al. 1972) Baer et al. 2000. <== *Bdellovibrio starrii* (basonym). *Peredibacter starrii* (new combination). Synonym links through N4Lid: 10.1601/ex.3681. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 15145. High-quality 16S rRNA sequence S000428666 (RDP), AF084852 (Genbank). N4Lid DOI: 10.1601/nm.3685

† *Bdellovibrio starrii*^{AL} Seidler et al. 1972. ==> *Bacteriovorax starrii* (new combination). Synonym links through N4Lid: 10.1601/ex.3681. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 15145. High-quality 16S rRNA sequence S000428666 (RDP), AF084852 (Genbank). N4Lid DOI: 10.1601/nm.3681

Order Myxococcales^{AL} Tchan et al 1948. N4Lid DOI: 10.1601/nm.3690

Suborder "Cystobacterineae" N4Lid DOI: 10.1601/nm.3691

Family Cystobacteraceae^{AL} McCurdy 1970. N4Lid DOI: 10.1601/nm.3692

Genus Cystobacter^{AL} Schroeter 1886. N4Lid DOI: 10.1601/nm.3693

Cystobacter fuscus^{AL} Schroeter 1886. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 25194. High-quality 16S rRNA sequence S000436919 (RDP), M94276 (Genbank). N4Lid DOI: 10.1601/nm.3694

"*Cystobacter armeniaca*" Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 14710. N4Lid DOI: 10.1601/nm.3695

"*Cystobacter badius*" Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 17723. N4Lid DOI: 10.1601/nm.3696

"*Cystobacter disciformis*" <==Synonym links through N4Lid: 10.1601/ex.3697. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 33172. N4Lid DOI: 10.1601/nm.3697

† *Angiococcus disciformis*^{VP} (Thaxter 1904) Hook et al. 1980. <== *Myxococcus disciformis* (basonym). Synonym links through N4Lid: 10.1601/ex.3697. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 33172. N4Lid DOI: 10.1601/nm.9089

† *Myxococcus disciformis*^{AL} Thaxter 1904. ==> *Angiococcus disciformis* (new combination). Synonym links through N4Lid: 10.1601/ex.3697. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 33172. N4Lid DOI: 10.1601/nm.3722

Cystobacter ferrugineus^{AL} (Krzemieniewska and Krzemieniewski 1927) McCurdy 1970. Source of type material recommended for DOE sponsored genome sequencing by the JGI: Windsor M-203. High-quality 16S rRNA sequence S000539278 (RDP), AJ233901 (Genbank). N4Lid DOI: 10.1601/nm.3698

"*Cystobacter gracilis*" Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 14753. N4Lid DOI: 10.1601/nm.3699

"*Cystobacter miniatus*" Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 14712. N4Lid DOI: 10.1601/nm.3700

"*Cystobacter minor*" Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 14751. N4Lid DOI: 10.1601/nm.3701

"*Cystobacter velatus*" Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 14718. N4Lid DOI: 10.1601/nm.3702

"*Cystobacter violaceus*" Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 14727. N4Lid DOI: 10.1601/nm.3703

Genus Anaeromyxobacter^{VP} Sanford et al. 2002. GOLD ID: Gi01322. GCAT ID: 001731_GCAT. Entrez genome id: 17729. Sequenced strain: Fw109-5 is from a non-type strain. Genome sequencing is incomplete. Number of genomes of this species sequenced 2 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3704

Anaeromyxobacter dehalogenans^{VP} Sanford et al. 2002. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-258. High-quality 16S rRNA sequence S000392790 (RDP), AF382396 (Genbank). GOLD ID: Gc00340. GCAT ID: 000270_GCAT. Entrez genome id: 12634. Sequenced strain: 2CP-C is from a non-type strain. Genome sequencing is completed. Number of genomes of this species sequenced 2 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3705

Genus *Archangium*^{AL} Jahn 1924. N4Lid DOI: 10.1601/nm.3706

Archangium gephyra^{AL} Jahn 1924. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 25201. High-quality 16S rRNA sequence S000436916 (RDP), M94273 (Genbank). N4Lid DOI: 10.1601/nm.3707

Genus "*Hyalangium*" N4Lid DOI: 10.1601/nm.3708

"Hyalangium minutum" Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 14724. N4Lid DOI: 10.1601/nm.3709

Genus *Melittangium*^{AL} Jahn 1924. N4Lid DOI: 10.1601/nm.3710

Melittangium boletus^{AL} Jahn 1924. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 14713. High-quality 16S rRNA sequence S000539285 (RDP), AJ233908 (Genbank). N4Lid DOI: 10.1601/nm.3711

Melittangium alboraceum^{AL} (Peterson 1959) McCurdy 1971. Source of type material recommended for DOE sponsored genome sequencing by the JGI: No viable type strain.. N4Lid DOI: 10.1601/nm.3712

Melittangium lichenicola^{AL} (Thaxter 1892) McCurdy 1971. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 25944. High-quality 16S rRNA sequence S000436920 (RDP), M94277 (Genbank). N4Lid DOI: 10.1601/nm.3713

Genus *Stigmatella*^{AL} Berkeley and Curtis 1875. N4Lid DOI: 10.1601/nm.3714

Stigmatella aurantiaca^{AL} Berkeley and Curtis 1875. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 25190. High-quality 16S rRNA sequence S000436924 (RDP), M94281 (Genbank). GOLD ID: Gi00604. GCAT ID: 001526_GCAT. Sequenced strain: DW4 /3-1 is from a non-type strain. Genome sequencing is incomplete. Number of genomes of this species sequenced 2 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3715

Stigmatella erecta^{AL} (Schroeter 1886) McCurdy 1971. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 25191. High-quality 16S rRNA sequence S000540099 (RDP), AJ970180 (Genbank). N4Lid DOI: 10.1601/nm.3716

"Stigmatella hybrida" Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 14722. N4Lid DOI: 10.1601/nm.3717

Family *Myxococcaceae*^{AL} Jahn 1924. N4Lid DOI: 10.1601/nm.3718

Genus *Myxococcus*^{AL} Thaxter 1892. N4Lid DOI: 10.1601/nm.3719

Myxococcus fulvus^{AL} (Cohn 1875) Jahn 1911. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 25199. N4Lid DOI: 10.1601/nm.3720

Myxococcus flavescens^{VP} Yamanaka et al. 1990. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 51243. N4Lid DOI: 10.1601/nm.3723

Myxococcus macrosporus^{AL} (Krzemieniewska and Krzemieniewski 1926) Zahler and McCurdy 1974. Source of type material recommended for DOE sponsored genome sequencing by the JGI: Windsor M271. N4Lid DOI: 10.1601/nm.3724

Myxococcus stipitatus^{AL} Thaxter 1897. Source of type material recommended for DOE sponsored genome sequencing by the JGI: Windsor M78. N4Lid DOI: 10.1601/nm.3725

Myxococcus virescens^{AL} Thaxter 1892. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 25203. N4Lid DOI: 10.1601/nm.3726

Myxococcus xanthus^{AL} Beebe 1941. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 25232. High-quality 16S rRNA sequence S000750061 (RDP). GOLD ID: Gc00391. GCAT ID: 000219_GCAT. Entrez genome id: 1421. Sequenced strain: DK 1622 is from a non-type strain. Genome sequencing is completed. Number of genomes of this species sequenced 2 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3727

Genus † *Angiococcus*^{VP} (ex Jahn 1924) Hook et al. 1980. Unified with *Myxococcus*, links through 10.1601/nm.3719. N4Lid 10.1601/nm.9088

Genus "*Corallococcus*" N4Lid DOI: 10.1601/nm.3728

"Corallococcus coralloides" <==Synonym links through N4Lid: 10.1601/ex.3721. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 25202. High-quality 16S rRNA sequence S000367196 (RDP), AJ811588 (Genbank). N4Lid DOI: 10.1601/nm.3729

† *Myxococcus coralloides*^{AL} Thaxter 1892. ==> *Corallocooccus coralloides* (new combination).
Synonym links through N4Lid: 10.1601/ex.3721. Source of type material recommended for DOE
sponsored genome sequencing by the JGI: ATCC 25202. High-quality 16S rRNA sequence
S000367196 (RDP), AJ811588 (Genbank). N4Lid DOI: 10.1601/nm.3721

"*Corallocooccus exiguus*" Source of type material recommended for DOE sponsored genome
sequencing by the JGI: DSM 14696. High-quality 16S rRNA sequence S000367206 (RDP),
AJ811598 (Genbank). N4Lid DOI: 10.1601/nm.3730

"*Corallocooccus macrosporus*" Source of type material recommended for DOE sponsored genome
sequencing by the JGI: DSM 14697. N4Lid DOI: 10.1601/nm.3731

Genus "*Pyxicoccus*" N4Lid DOI: 10.1601/nm.3732

"*Pyxicoccus fallax*" Source of type material recommended for DOE sponsored genome
sequencing by the JGI: DSM 14698. N4Lid DOI: 10.1601/nm.3733

Suborder "*Sorangineae*" N4Lid DOI: 10.1601/nm.3734

Family *Polyangiaceae*^{AL} Jahn 1924. N4Lid DOI: 10.1601/nm.3735

Genus *Polyangium*^{AL} Link 1809. N4Lid DOI: 10.1601/nm.3736

Polyangium vitellinum^{AL} Link 1809. Source of type material recommended for DOE sponsored
genome sequencing by the JGI: TC 4564. N4Lid DOI: 10.1601/nm.3737

Polyangium aureum^{VP} (ex Krzemieniewska and Krzemieniewski 1930) Brockman 1989. Source
of type material recommended for DOE sponsored genome sequencing by the JGI: No viable type
strain.. N4Lid DOI: 10.1601/nm.3738

Polyangium fumosum^{AL} Krzemieniewska and Krzemieniewski 1930. Source of type material
recommended for DOE sponsored genome sequencing by the JGI: Windsor M257. N4Lid DOI:
10.1601/nm.3740

Polyangium luteum^{VP} (ex Krzemieniewska and Krzemieniewski 1927) Brockman 1989. Source
of type material recommended for DOE sponsored genome sequencing by the JGI: No viable type
strain.. N4Lid DOI: 10.1601/nm.3741

Polyangium parasiticum^{VP} (ex Geitler 1924) Brockman 1989. Source of type material
recommended for DOE sponsored genome sequencing by the JGI: No viable type strain.. N4Lid
DOI: 10.1601/nm.3743

Polyangium solediatum^{VP} (ex Thaxter 1904) Brockman 1989. Source of type material
recommended for DOE sponsored genome sequencing by the JGI: DSM 14670. N4Lid DOI:
10.1601/nm.3745

Polyangium spumosum^{VP} (ex Krzemieniewska and Krzemieniewski 1927) Brockman 1989.
Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM
14734. N4Lid DOI: 10.1601/nm.3746

Genus "*Byssovorax*" N4Lid DOI: 10.1601/nm.3747

"*Byssovorax cruenta*" Source of type material recommended for DOE sponsored genome
sequencing by the JGI: DSM 14553. High-quality 16S rRNA sequence S000417333 (RDP),
AJ833647 (Genbank). N4Lid DOI: 10.1601/nm.3748

Genus *Chondromyces*^{AL} Berkeley and Curtis 1874. N4Lid DOI: 10.1601/nm.3749

Chondromyces crocatus^{AL} Berkeley and Curtis 1874. Source of type material recommended for
DOE sponsored genome sequencing by the JGI: DSM 14714. N4Lid DOI: 10.1601/nm.3750

Chondromyces apiculatus^{AL} Thaxter 1897. Source of type material recommended for DOE
sponsored genome sequencing by the JGI: DSM 14605. High-quality 16S rRNA sequence
S000539315 (RDP), AJ233938 (Genbank). N4Lid DOI: 10.1601/nm.3751

Chondromyces catenulatus^{AL} Thaxter 1904. Source of type material recommended for DOE
sponsored genome sequencing by the JGI: TC 4517. N4Lid DOI: 10.1601/nm.3752

Chondromyces lanuginosus^{AL} Kofler 1913. Source of type material recommended for DOE
sponsored genome sequencing by the JGI: DSM 14631. High-quality 16S rRNA sequence
S000539316 (RDP), AJ233939 (Genbank). N4Lid DOI: 10.1601/nm.3753

Chondromyces pediculatus^{AL} Thaxter 1904. Source of type material recommended for DOE
sponsored genome sequencing by the JGI: DSM 14607. N4Lid DOI: 10.1601/nm.3754

"Chondromyces robustus" Source of type material recommended for DOE sponsored genome sequencing by the JGI: No viable type strain.. High-quality 16S rRNA sequence S000539319 (RDP), AJ233942 (Genbank). N4Lid DOI: 10.1601/nm.3755

Genus "*Haploangium*" N4Lid DOI: 10.1601/nm.3756

"Haploangium rugiseptum" <==Synonym links through N4Lid: 10.1601/ex.3744. Source of type material recommended for DOE sponsored genome sequencing by the JGI: UMH Peterson 51. N4Lid DOI: 10.1601/nm.3757

† *Polyangium rugiseptum*^{AL} (Peterson 1959) McCurdy 1970. ==> *Haploangium rugiseptum* (new combination). Synonym links through N4Lid: 10.1601/ex.3744. Source of type material recommended for DOE sponsored genome sequencing by the JGI: UMH Peterson 51. N4Lid DOI: 10.1601/nm.3744

"Haploangium minus" <==Synonym links through N4Lid: 10.1601/ex.3742. Source of type material recommended for DOE sponsored genome sequencing by the JGI: UMH Peterson 41. N4Lid DOI: 10.1601/nm.3758

† *Polyangium minor*^{AL} (Peterson 1959) McCurdy 1970. ==> *Haploangium minus* (new combination). Synonym links through N4Lid: 10.1601/ex.3742. Source of type material recommended for DOE sponsored genome sequencing by the JGI: UMH Peterson 41. N4Lid DOI: 10.1601/nm.3742

Genus "*Jahnia*" N4Lid DOI: 10.1601/nm.3759

"Jahnia thaxteri" Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 14626. N4Lid DOI: 10.1601/nm.3760

Genus "*Sorangium*" N4Lid DOI: 10.1601/nm.3761

"Sorangium cellulosum" <==Synonym links through N4Lid: 10.1601/ex.3739. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 14627. GOLD ID: Gi00561. GCAT ID: 001504_GCAT. Sequenced strain: So ce56 is from a non-type strain. Genome sequencing is incomplete. Number of genomes of this species sequenced 1 (GOLD). N4Lid DOI: 10.1601/nm.3762

† *Polyangium cellulosum*^{VP} (ex Imshenetski and Solntseva 1936) Brockman 1989. ==> *Sorangium cellulosum* (new combination). Synonym links through N4Lid: 10.1601/ex.3739. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 14627. GOLD ID: Gi00561. GCAT ID: 001504_GCAT. Sequenced strain: So ce56 is from a non-type strain. Genome sequencing is incomplete. Number of genomes of this species sequenced 1 (GOLD) 0 (NCBI). N4Lid DOI: 10.1601/nm.3739

Suborder "*Nannocystineae*" N4Lid DOI: 10.1601/nm.3763

Family *Nannocystaceae*^{VP} Reichenbach 2006. N4Lid DOI: 10.1601/nm.3764

Genus *Nannocystis*^{AL} Reichenbach 1970. N4Lid DOI: 10.1601/nm.3765

"Nannocystis excedens subsp. excedans" Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 25963. N4Lid DOI: 10.1601/nm.3766

"Nannocystis excedens subsp. aggregans" Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 25963. N4Lid DOI: 10.1601/nm.3767

"Nannocystis excedens subsp. cinnabarina" Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 14641. N4Lid DOI: 10.1601/nm.3768

"Nannocystis excedens subsp. glomerata" Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 14640. N4Lid DOI: 10.1601/nm.3769

"Nannocystis excedens subsp. pulla" Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 14629. N4Lid DOI: 10.1601/nm.3770

"Nannocystis pusilla" Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 14622. N4Lid DOI: 10.1601/nm.3771

Genus *Enhygromyxa*^{VP} Iizuka et al. 2003. N4Lid DOI: 10.1601/nm.3772

Enhygromyxa salina^{VP} Iizuka et al. 2003. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 15217. High-quality 16S rRNA sequence S000385672 (RDP), AB097590 (Genbank). N4Lid DOI: 10.1601/nm.3773

Genus *Plesiocystis*^{VP} Iizuka et al. 2003. N4Lid DOI: 10.1601/nm.3774

Plesiocystis pacifica^{VP} Iizuka et al. 2003. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 14875. High-quality 16S rRNA sequence S000384935 (RDP), AB083432 (Genbank). GOLD ID: Gi01444. GCAT ID: 000909_GCAT. Sequenced strain: SIR-1 is type material. Genome sequencing is incomplete. Number of genomes of this species sequenced 1 (GOLD). N4Lid DOI: 10.1601/nm.3775

Family "Haliangiaceae" N4Lid DOI: 10.1601/nm.3776

Genus *Haliangium*^{VP} Fudou et al. 2002. N4Lid DOI: 10.1601/nm.3777

Haliangium ochraceum^{VP} Fudou et al. 2002. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 14365. High-quality 16S rRNA sequence S000382370 (RDP), AB016470 (Genbank). N4Lid DOI: 10.1601/nm.3778

Haliangium tepidum^{VP} Fudou et al. 2002. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 14436. High-quality 16S rRNA sequence S000022588 (RDP), AB062751 (Genbank). N4Lid DOI: 10.1601/nm.3779

Family "Kofleriaceae" N4Lid DOI: 10.1601/nm.3780

Genus "Kofleria" N4Lid DOI: 10.1601/nm.3781

"*Kofleria flava*" Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 14601. N4Lid DOI: 10.1601/nm.3782

Class Epsilonproteobacteria^{VP} Garrity et al 2006. N4Lid DOI: 10.1601/nm.3783

Order Campylobacterales^{VP} Garrity et al 2006. N4Lid DOI: 10.1601/nm.3784

Family Campylobacteraceae^{VP} Vandamme and De Ley 1991. N4Lid DOI: 10.1601/nm.3785

Genus *Campylobacter*^{AL} Sebald and Véron 1963. N4Lid DOI: 10.1601/nm.3786

Campylobacter fetus subsp. fetus^{AL} (Smith and Taylor 1919) Véron and Chatelain 1973. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 19438. High-quality 16S rRNA sequence S000414266 (RDP), L04314 (Genbank). GOLD ID: Gc00466. GCAT ID: 000143_GCAT. Entrez genome id: 16293. Sequenced strain: 82-40 is from a non-type strain. Genome sequencing is completed. Number of genomes of this species sequenced 2 (GOLD) 2 (NCBI). N4Lid DOI: 10.1601/nm.3787

Campylobacter fetus subsp. venerealis^{AL} (Florent 1959) Véron and Chatelain 1973. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 19438. N4Lid DOI: 10.1601/nm.3795

Campylobacter coli^{AL} (Doyle 1948) Véron and Chatelain 1973. ≈ *Campylobacter hyoilei* (later heterotypic synonym), links through 10.1601/nm.3790. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 33559. High-quality 16S rRNA sequence S000436447 (RDP), M59073 (Genbank). GOLD ID: Gi00118. GCAT ID: 000819_GCAT. Entrez genome id: 12516. Sequenced strain: ATCC BAA-1061 is from a non-type strain. Genome sequencing is incomplete. Number of genomes of this species sequenced 1 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3790

† *Campylobacter hyoilei*^{VP} Alderton et al. 1995. ≈ *Campylobacter coli* (earlier heterotypic synonym), links through 10.1601/nm.3790. Source of type material recommended for DOE sponsored genome sequencing by the JGI: CCUG 33450. N4Lid DOI: 10.1601/nm.3799

Campylobacter concisus^{VP} Tanner et al. 1981. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 33237. High-quality 16S rRNA sequence S000414274 (RDP), L04322 (Genbank). GOLD ID: Gi01266. GCAT ID: 000829_GCAT. Entrez genome id: 17159. Sequenced strain: 13826 is from a non-type strain. Genome sequencing is incomplete. Number of genomes of this species sequenced 1 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3791

Campylobacter curvus^{VP} (Tanner et al. 1984) Vandamme et al. 1991. <== *Wolinella curva* (basonym). Synonym links through N4Lid: 10.1601/ex.3862. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 35224. High-quality 16S rRNA sequence S000414265 (RDP), L04313 (Genbank). GOLD ID: Gi01267. GCAT ID: 000830_GCAT. Entrez genome id: 17161. Sequenced strain: 526 is from a non-type strain. Genome sequencing is incomplete. Number of genomes of this species sequenced 1 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3793

† *Wolinella curva*^{VP} Tanner et al. 1984. ==> *Campylobacter curvus* (new combination). Synonym links through N4Lid: 10.1601/ex.3862. Source of type material recommended for DOE

sponsored genome sequencing by the JGI: ATCC 35224. High-quality 16S rRNA sequence S000414265 (RDP), L04313 (Genbank). GOLD ID: Gi01267. GCAT ID: 000830_GCAT. Entrez genome id: 17161. Sequenced strain: 526 is from a non-type strain. Genome sequencing is incomplete. Number of genomes of this species sequenced 1 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3862

Campylobacter gracilis^{VP} (Tanner et al. 1981) Vandamme et al. 1995. <== *Bacteroides gracilis* (**basonym**). Synonym links through N4Lid: 10.1601/ex.7954. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 33236. High-quality 16S rRNA sequence S000414272 (RDP), L04320 (Genbank). N4Lid DOI: 10.1601/nm.3796

† *Bacteroides gracilis*^{VP} Tanner et al. 1981. ==> *Campylobacter gracilis* (**new combination**). Synonym links through N4Lid: 10.1601/ex.7954. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 33236. High-quality 16S rRNA sequence S000414272 (RDP), L04320 (Genbank). N4Lid DOI: 10.1601/nm.7954

Campylobacter helveticus^{VP} Stanley et al. 1993. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 51209. High-quality 16S rRNA sequence S000437001 (RDP), U03022 (Genbank). N4Lid DOI: 10.1601/nm.3797

Campylobacter hominis^{VP} Lawson et al. 2001. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-381. High-quality 16S rRNA sequence S000004853 (RDP), AJ251584 (Genbank). GOLD ID: Gi01268. GCAT ID: 000831_GCAT. Sequenced strain: LMG 19568 is type material. Genome sequencing is incomplete. Number of genomes of this species sequenced 1 (GOLD). N4Lid DOI: 10.1601/nm.3798

Campylobacter hyointestinalis subsp. hyointestinalis^{VP} Gebhart et al. 1985. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 35217. High-quality 16S rRNA sequence S000436628 (RDP), M65010 (Genbank). N4Lid DOI: 10.1601/nm.3800

Campylobacter hyointestinalis subsp. lawsonii^{VP} On et al. 1995. Source of type material recommended for DOE sponsored genome sequencing by the JGI: CCUG 34538. High-quality 16S rRNA sequence S000428836 (RDP), AF097685 (Genbank). N4Lid DOI: 10.1601/nm.3801

Campylobacter insulaenigrae^{VP} Foster et al. 2004. Source of type material recommended for DOE sponsored genome sequencing by the JGI: CCUG 48653. High-quality 16S rRNA sequence S000440230 (RDP), AJ620504 (Genbank). N4Lid DOI: 10.1601/nm.8943

Campylobacter jejuni subsp. jejuni^{VP} (Jones et al. 1931) Véron and Chatelain 1973. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 33560. High-quality 16S rRNA sequence S000414431 (RDP), L14630 (Genbank). GOLD ID: Gc00025. GCAT ID: 000585_GCAT. Entrez genome id: 8. Sequenced strain: NCTC 11168 is from a non-type strain. Genome sequencing is completed. Number of genomes of this species sequenced 10 (GOLD) 10 (NCBI). N4Lid DOI: 10.1601/nm.3802

Campylobacter jejuni subsp. doylei^{VP} Steele and Owen 1988. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 49349. High-quality 16S rRNA sequence S000690671 (RDP). GOLD ID: Gi01293. GCAT ID: 000828_GCAT. Entrez genome id: 17163. Sequenced strain: 270 is from a non-type strain. Genome sequencing is incomplete. Number of genomes of this species sequenced 1 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3803

Campylobacter lanienae^{VP} Logan et al. 2000. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-633. High-quality 16S rRNA sequence S000428017 (RDP), AF043425 (Genbank). N4Lid DOI: 10.1601/nm.3804

Campylobacter lari^{VP} corrig. Benjamin et al. 1984. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 35221. High-quality 16S rRNA sequence S000414268 (RDP), L04316 (Genbank). GOLD ID: Gi00122. GCAT ID: 000832_GCAT. Entrez genome id: 12517. Sequenced strain: ATCC BAA-1060 is from a non-type strain. Genome sequencing is incomplete. Number of genomes of this species sequenced 1 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3805

Campylobacter mucosalis^{VP} (Lawson et al. 1981) Roop et al. 1985. <== *Campylobacter sputorum subsp. mucosalis* (**basonym**). Synonym links through N4Lid: 10.1601/ex.3815. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 43264. N4Lid DOI: 10.1601/nm.3806

† *Campylobacter sputorum subsp. mucosalis*^{VP} (ex Lawson and Rowland 1974) Lawson et al. 1981. ==> *Campylobacter mucosalis* (new combination). Synonym links through N4Lid: 10.1601/ex.3815. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 43264. N4Lid DOI: 10.1601/nm.3815

Campylobacter rectus^{VP} (Tanner et al. 1981) Vandamme et al. 1991. <== *Wolinella recta* (basonym). Synonym links through N4Lid: 10.1601/ex.3863. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 33238. High-quality 16S rRNA sequence S000414269 (RDP), L04317 (Genbank). N4Lid DOI: 10.1601/nm.3811

† *Wolinella recta*^{VP} Tanner et al. 1981. ==> *Campylobacter rectus* (new combination). Synonym links through N4Lid: 10.1601/ex.3863. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 33238. High-quality 16S rRNA sequence S000414269 (RDP), L04317 (Genbank). N4Lid DOI: 10.1601/nm.3863

Campylobacter showae^{VP} Etoh et al. 1993. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 51146. High-quality 16S rRNA sequence S000414317 (RDP), L06974 (Genbank). N4Lid DOI: 10.1601/nm.3812

Campylobacter sputorum subsp. sputorum^{AL} (Prévot 1940) Véron and Chatelain 1973. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 33562. High-quality 16S rRNA sequence S000021698 (RDP), X67775 (Genbank). N4Lid DOI: 10.1601/nm.3813

Campylobacter sputorum subsp. bubulus^{AL} (Florent 1953) Véron and Chatelain 1973. Source of type material recommended for DOE sponsored genome sequencing by the JGI: CCUG 11289. N4Lid DOI: 10.1601/nm.3814

Campylobacter upsaliensis^{VP} Sandstedt and Ursing 1991. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 43954. High-quality 16S rRNA sequence S000414429 (RDP), L14628 (Genbank). GOLD ID: Gi00123. GCAT ID: 000833_GCAT. Entrez genome id: 12518. Sequenced strain: ATCC BAA-1059 is from a non-type strain. Genome sequencing is incomplete. Number of genomes of this species sequenced 1 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3816

Genus *Arcobacter*^{VP} Vandamme et al. 1991. N4Lid DOI: 10.1601/nm.3817

Arcobacter nitrofigilis^{VP} (McClung et al. 1983) Vandamme et al. 1991. <== *Campylobacter nitrofigilis* (basonym). Synonym links through N4Lid: 10.1601/ex.3808. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 33309. High-quality 16S rRNA sequence S000414428 (RDP), L14627 (Genbank). N4Lid DOI: 10.1601/nm.3818

† *Campylobacter nitrofigilis*^{VP} McClung et al. 1983. ==> *Arcobacter nitrofigilis* (new combination). Synonym links through N4Lid: 10.1601/ex.3808. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 33309. High-quality 16S rRNA sequence S000414428 (RDP), L14627 (Genbank). N4Lid DOI: 10.1601/nm.3808

Arcobacter butzleri^{VP} (Kiehlbauch et al. 1991) Vandamme et al. 1992. <== *Campylobacter butzleri* (basonym). Synonym links through N4Lid: 10.1601/ex.3788. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 49616. N4Lid DOI: 10.1601/nm.3819

† *Campylobacter butzleri*^{VP} Kiehlbauch et al. 1991. ==> *Arcobacter butzleri* (new combination). Synonym links through N4Lid: 10.1601/ex.3788. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 49616. N4Lid DOI: 10.1601/nm.3788

Arcobacter cibarius^{VP} Houf et al. 2005. Source of type material recommended for DOE sponsored genome sequencing by the JGI: CCUG 48482. High-quality 16S rRNA sequence S000471105 (RDP), AJ607391 (Genbank). N4Lid 10.1601/nm.9357

Arcobacter cryaerophilus^{VP} (Neill et al. 1985) Vandamme et al. 1991. <== *Campylobacter cryaerophilus* (basonym). Synonym links through N4Lid: 10.1601/ex.3792. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 43158. High-quality 16S rRNA sequence S000414425 (RDP), L14624 (Genbank). N4Lid DOI: 10.1601/nm.3820

† *Campylobacter cryaerophilus*^{VP} corrig. Neill et al. 1985. ==> *Arcobacter cryaerophilus* (new combination). Synonym links through N4Lid: 10.1601/ex.3792. Source of type material

recommended for DOE sponsored genome sequencing by the JGI: ATCC 43158. High-quality 16S rRNA sequence S000414425 (RDP), L14624 (Genbank). N4Lid DOI: 10.1601/nm.3792

Arcobacter halophilus^{VP} Donachie et al. 2005. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-1022. High-quality 16S rRNA sequence S000543775 (RDP), AF513455 (Genbank). N4Lid 10.1601/nm.9451

Arcobacter skirrowii^{VP} Vandamme et al. 1992. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 51132. High-quality 16S rRNA sequence S000414426 (RDP), L14625 (Genbank). N4Lid DOI: 10.1601/nm.3821

Genus † *Dehalospirillum*^{VP} Scholz-Muramatsu et al. 2002. ≈ *Sulfurospirillum* (earlier heterotypic synonym), links through 10.1601./nm.3824. N4Lid DOI: 10.1601/nm.3822

Genus *Sulfurospirillum*^{VP} Schumacher et al. 1993. ≈ *Sulfurospirillum* (later heterotypic synonym), links through 10.1601./nm.3824. N4Lid DOI: 10.1601/nm.3824

Sulfurospirillum deleyianum^{VP} Schumacher et al. 1993. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 51133. High-quality 16S rRNA sequence S000128537 (RDP), Y13671 (Genbank). N4Lid DOI: 10.1601/nm.3825

Sulfurospirillum arcachonense^{VP} Finster et al. 1997. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 700423. High-quality 16S rRNA sequence S000018038 (RDP), Y11561 (Genbank). N4Lid DOI: 10.1601/nm.3826

Sulfurospirillum arsenophilum^{VP} Stolz et al. 1999. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 700056. High-quality 16S rRNA sequence S000438625 (RDP), U85964 (Genbank). N4Lid DOI: 10.1601/nm.3827

Sulfurospirillum barnesii^{VP} Stolz et al. 1999. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 700032. High-quality 16S rRNA sequence S000427973 (RDP), AF038843 (Genbank). N4Lid DOI: 10.1601/nm.3828

Sulfurospirillum halorespirans^{VP} Luijten et al. 2003. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-583. High-quality 16S rRNA sequence S000389334 (RDP), AF218076 (Genbank). N4Lid DOI: 10.1601/nm.3829

Sulfurospirillum multivorans^{VP} (Scholz-Muramatsu et al. 2002) Luijten et al. 2003. <== *Dehalospirillum multivorans* (basonym). Synonym links through N4Lid: 10.1601/ex.3823. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 12446. N4Lid DOI: 10.1601/nm.3830

† *Dehalospirillum multivorans*^{VP} Scholz-Muramatsu et al. 2002. ==> *Sulfurospirillum multivorans* (new combination). Synonym links through N4Lid: 10.1601/ex.3823. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 12446. N4Lid DOI: 10.1601/nm.3823

Family *Helicobacteraceae*^{VP} Garrity et al 2006. N4Lid DOI: 10.1601/nm.3831

Genus *Helicobacter*^{VP} Goodwin et al. 1989. N4Lid DOI: 10.1601/nm.3832

Helicobacter pylori^{VP} (Marshall et al. 1985) Goodwin et al. 1989. <== *Campylobacter pylori subsp. pylori* (basonym). Synonym links through N4Lid: 10.1601/ex.3809. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 43504. High-quality 16S rRNA sequence S000014640 (RDP), Z25741 (Genbank). GOLD ID: Gc00007. GCAT ID: 000603_GCAT. Entrez genome id: 233. Sequenced strain: ATCC 700392 is from a non-type strain. Genome sequencing is completed. Number of genomes of this species sequenced 6 (GOLD) 3 (NCBI). N4Lid DOI: 10.1601/nm.3833

† *Campylobacter pylori subsp. pylori*^{VP} Marshall et al. 1985. ==> *Helicobacter pylori* (new combination). Synonym links through N4Lid: 10.1601/ex.3809. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 43504. High-quality 16S rRNA sequence S000436866 (RDP), M88157 (Genbank). GOLD ID: Gc00007. GCAT ID: 000603_GCAT. Entrez genome id: 233. Sequenced strain: ATCC 700392 is from a non-type strain. Genome sequencing is completed. Number of genomes of this species sequenced 6 (GOLD) 3 (NCBI). N4Lid DOI: 10.1601/nm.3809

† *Helicobacter nemestrinae*^{VP} Bronsdon et al. 1991. ≈ *Helicobacter pylori* (earlier heterotypic synonym), links through 10.1601./nm.3833. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 49396. N4Lid DOI: 10.1601/nm.3849

- Helicobacter acinonychis*^{VP} corrig. Eaton et al. 1993. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 51101. High-quality 16S rRNA sequence S000363188 (RDP), M88148 (Genbank). GOLD ID: Gc00403. GCAT ID: 000212_GCAT. Entrez genome id: 17251. Sequenced strain: Sheeba is from a non-type strain. Genome sequencing is completed. Number of genomes of this species sequenced 1 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3834
- Helicobacter anseris*^{VP} Fox et al. 2006. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-1299. High-quality 16S rRNA sequence S000651603 (RDP), DQ415545 (Genbank). N4Lid DOI: 10.1601/nm.10315
- Helicobacter aurati*^{VP} Patterson et al. 2002. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-1. High-quality 16S rRNA sequence S000390922 (RDP), AF297868 (Genbank). N4Lid DOI: 10.1601/nm.3835
- Helicobacter bilis*^{VP} Fox et al. 1997. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 51630. High-quality 16S rRNA sequence S000363265 (RDP), U18766 (Genbank). N4Lid DOI: 10.1601/nm.3836
- Helicobacter bizzozeronii*^{VP} Hänninen et al. 1996. Source of type material recommended for DOE sponsored genome sequencing by the JGI: CCUG 35545. High-quality 16S rRNA sequence S000145295 (RDP), Y09404 (Genbank). N4Lid DOI: 10.1601/nm.3837
- Helicobacter brantae*^{VP} Fox et al. 2006. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-1298. High-quality 16S rRNA sequence S000651604 (RDP), DQ415546 (Genbank). N4Lid DOI: 10.1601/nm.10316
- Helicobacter canadensis*^{VP} Fox et al. 2002. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 700968. High-quality 16S rRNA sequence S000390281 (RDP), AF262037 (Genbank). N4Lid DOI: 10.1601/nm.3838
- Helicobacter canis*^{VP} Stanley et al. 1994. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 51401. High-quality 16S rRNA sequence S000414408 (RDP), L13464 (Genbank). N4Lid DOI: 10.1601/nm.3839
- Helicobacter cetorum*^{VP} Harper et al. 2006. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-540. High-quality 16S rRNA sequence S000390782 (RDP), AF292378 (Genbank). N4Lid DOI: 10.1601/nm.10317
- Helicobacter cholecystus*^{VP} Franklin et al. 1997. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 700242. High-quality 16S rRNA sequence S000437657 (RDP), U46129 (Genbank). N4Lid DOI: 10.1601/nm.3840
- Helicobacter cinaedi*^{VP} (Totten et al. 1988) Vandamme et al. 1991. <== *Campylobacter cinaedi* (basonym). Synonym links through N4Lid: 10.1601/ex.3789. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-847. High-quality 16S rRNA sequence S000334505 (RDP), M88150 (Genbank). N4Lid DOI: 10.1601/nm.3841
- † *Campylobacter cinaedi*^{VP} Totten et al. 1988. ==> *Helicobacter cinaedi* (new combination). Synonym links through N4Lid: 10.1601/ex.3789. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-847. High-quality 16S rRNA sequence S000334505 (RDP), M88150 (Genbank). N4Lid DOI: 10.1601/nm.3789
- Helicobacter cynogastricus*^{VP} Van den Bulck et al. 2006. Source of type material recommended for DOE sponsored genome sequencing by the JGI: LMG 23188. High-quality 16S rRNA sequence S000536639 (RDP), DQ004689 (Genbank). N4Lid DOI: 10.1601/nm.10159
- Helicobacter felis*^{VP} Paster et al. 1991. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 49179. High-quality 16S rRNA sequence S000375738 (RDP), M57398 (Genbank). N4Lid DOI: 10.1601/nm.3842
- Helicobacter fennelliae*^{VP} (Totten et al. 1988) Vandamme et al. 1991. <== *Campylobacter fennelliae* (basonym). Synonym links through N4Lid: 10.1601/ex.3794. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 35684. High-quality 16S rRNA sequence S000363190 (RDP), M88154 (Genbank). N4Lid DOI: 10.1601/nm.3843
- † *Campylobacter fennelliae*^{VP} Totten et al. 1988. ==> *Helicobacter fennelliae* (new combination). Synonym links through N4Lid: 10.1601/ex.3794. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 35684. High-quality 16S rRNA sequence S000363190 (RDP), M88154 (Genbank). N4Lid DOI: 10.1601/nm.3794

- Helicobacter ganmani*^{VP} Robertson et al. 2001. Source of type material recommended for DOE sponsored genome sequencing by the JGI: CCUG 43526. High-quality 16S rRNA sequence S000386802 (RDP), AF000221 (Genbank). N4Lid DOI: 10.1601/nm.3845
- "*Helicobacter heilmannii*" O'Rourke et al. 2004. Source of type material recommended for DOE sponsored genome sequencing by the JGI: No viable deposit available. High-quality 16S rRNA sequence S000014116 (RDP), AF506786 (Genbank). N4Lid 10.1601/nm.8973
- Helicobacter hepaticus*^{VP} Fox et al. 1994. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 51448. High-quality 16S rRNA sequence S000334525 (RDP), U07574 (Genbank). GOLD ID: Gc00139. GCAT ID: 000471_GCAT. Entrez genome id: 185. Sequenced strain: ATCC 51449 is from a non-type strain. Genome sequencing is completed. Number of genomes of this species sequenced 1 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3844
- Helicobacter marmotae*^{VP} Fox et al. 2006. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-546. High-quality 16S rRNA sequence S000391774 (RDP), AF333341 (Genbank). N4Lid DOI: 10.1601/nm.10318
- Helicobacter mastomyrinus*^{VP} Shen et al. 2006. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-1046. High-quality 16S rRNA sequence S000505602 (RDP), AY742307 (Genbank). N4Lid DOI: 10.1601/nm.10319
- Helicobacter mesocricetorum*^{VP} Simmons et al. 2000. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 700932. High-quality 16S rRNA sequence S000428427 (RDP), AF072471 (Genbank). N4Lid DOI: 10.1601/nm.3846
- Helicobacter muridarum*^{VP} Lee et al. 1992. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 49282. High-quality 16S rRNA sequence S000334502 (RDP), M80205 (Genbank). N4Lid DOI: 10.1601/nm.3847
- Helicobacter mustelae*^{VP} (Fox et al. 1988) Goodwin et al. 1989. <== *Campylobacter pylori* subsp. *mustelae* (basonym). Synonym links through N4Lid: 10.1601/ex.3810. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 43772. High-quality 16S rRNA sequence S000363169 (RDP), M35048 (Genbank). GOLD ID: Gi00294. GCAT ID: 001133_GCAT. Entrez genome id: 13426. Sequenced strain: ATCC 43772 is type material. Genome sequencing is incomplete. Number of genomes of this species sequenced 1 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3848
- † *Campylobacter mustelae*^{VP} (Fox et al. 1988) Fox et al. 1989. <== *Campylobacter pylori* subsp. *mustelae* (basonym). *Helicobacter mustelae* (new combination). Synonym links through N4Lid: 10.1601/ex.3810. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 43772. High-quality 16S rRNA sequence S000363169 (RDP), M35048 (Genbank). GOLD ID: Gi00294. GCAT ID: 001133_GCAT. Entrez genome id: 13426. Sequenced strain: ATCC 43772 is type material. Genome sequencing is incomplete. Number of genomes of this species sequenced 1 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3807
- † *Campylobacter pylori* subsp. *mustelae*^{VP} Fox et al. 1988. ==> *Campylobacter mustelae* (new combination). Synonym links through N4Lid: 10.1601/ex.3810. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 43772. High-quality 16S rRNA sequence S000363169 (RDP), M35048 (Genbank). GOLD ID: Gi00294. GCAT ID: 001133_GCAT. Entrez genome id: 13426. Sequenced strain: ATCC 43772 is type material. Genome sequencing is incomplete. Number of genomes of this species sequenced 1 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3810
- Helicobacter pametensis*^{VP} Dewhirst et al. 1994. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 51478. High-quality 16S rRNA sequence S000363187 (RDP), M88147 (Genbank). N4Lid DOI: 10.1601/nm.3850
- Helicobacter pullorum*^{VP} Stanley et al. 1995. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 51801. High-quality 16S rRNA sequence S000622336 (RDP), L36141 (Genbank). N4Lid DOI: 10.1601/nm.3851
- Helicobacter rodentium*^{VP} Shen et al. 1997. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 700285. High-quality 16S rRNA sequence S000438897 (RDP), U96296 (Genbank). N4Lid DOI: 10.1601/nm.3852
- Helicobacter salomonis*^{VP} Jalava et al. 1997. Source of type material recommended for DOE sponsored genome sequencing by the JGI: CCUG 37845. High-quality 16S rRNA sequence S000438741 (RDP), U89351 (Genbank). N4Lid DOI: 10.1601/nm.3853

Helicobacter trogontum^{VP} Mendes et al. 1996. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 700114. High-quality 16S rRNA sequence S000438115 (RDP), U65103 (Genbank). N4Lid DOI: 10.1601/nm.3854

Helicobacter typhlonius^{VP} Franklin et al. 2002. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-367. High-quality 16S rRNA sequence S000387334 (RDP), AF127912 (Genbank). N4Lid DOI: 10.1601/nm.3855

Genus *Sulfuricurvum*^{VP} Kodama and Watanabe 2004. N4Lid 10.1601/nm.8871

Sulfuricurvum kujiense^{VP} Kodama and Watanabe 2004. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-921. High-quality 16S rRNA sequence S000383585 (RDP), AB053951 (Genbank). N4Lid 10.1601/nm.8872

Genus *Sulfurimonas*^{VP} Inagaki et al. 2003. N4Lid DOI: 10.1601/nm.3856

Sulfurimonas autotrophica^{VP} Inagaki et al. 2003. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-671. High-quality 16S rRNA sequence S000252003 (RDP), AB088431 (Genbank). N4Lid DOI: 10.1601/nm.3857

Sulfurimonas denitrificans^{VP} (Timmer-ten Hoor 1975) Takai et al. 2006. <== *Thiomicrospira denitrificans* (basonym). Synonym links through N4Lid: 10.1601/ex.2331. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 33889. High-quality 16S rRNA sequence S000435956 (RDP), L40808 (Genbank). GOLD ID: Gc00328. GCAT ID: 000282_GCAT. Entrez genome id: 13019. Sequenced strain: ATCC 33889 is type material. Genome sequencing is completed. Number of genomes of this species sequenced 1 (GOLD) 1 (NCBI). N4Lid 10.1601/nm.9790

† *Thiomicrospira denitrificans*^{AL} Timmer-ten Hoor 1975. ==> *Sulfurimonas denitrificans* (new combination). Synonym links through N4Lid: 10.1601/ex.2331. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 33889. High-quality 16S rRNA sequence S000435956 (RDP), L40808 (Genbank). GOLD ID: Gc00328. GCAT ID: 000282_GCAT. Entrez genome id: 13019. Sequenced strain: ATCC 33889 is type material. Genome sequencing is completed. Number of genomes of this species sequenced 1 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.2331

Sulfurimonas parvalvinellae^{VP} Takai et al. 2006. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 17229. High-quality 16S rRNA sequence S000650635 (RDP), AB252048 (Genbank). N4Lid DOI: 10.1601/nm.10230

Genus *Sulfurovum*^{VP} Inagaki et al. 2004. N4Lid 10.1601/nm.8873

Sulfurovum lithotrophicum^{VP} Inagaki et al. 2004. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-797. High-quality 16S rRNA sequence S000415023 (RDP), AB091292 (Genbank). N4Lid 10.1601/nm.8874

Genus *Thiovulum*^{AL} Hinze 1913. N4Lid DOI: 10.1601/nm.3858

Thiovulum majus^{AL} Hinze 1913. Source of type material recommended for DOE sponsored genome sequencing by the JGI: no culture isolated. N4Lid DOI: 10.1601/nm.3859

Genus *Wolinella*^{VP} Tanner et al. 1981. N4Lid DOI: 10.1601/nm.3860

Wolinella succinogenes^{VP} (Wolin et al. 1961) Tanner et al. 1981. <== *Vibrio succinogenes* (basonym). Synonym links through N4Lid: 10.1601/ex.3006. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 29543. High-quality 16S rRNA sequence S000363191 (RDP), M88159 (Genbank). GOLD ID: Gc00158. GCAT ID: 000452_GCAT. Entrez genome id: 445. Sequenced strain: DSM 1740 is type material. Genome sequencing is completed. Number of genomes of this species sequenced 1 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3861

† *Vibrio succinogenes*^{AL} Wolin et al. 1961. ==> *Wolinella succinogenes* (new combination). Synonym links through N4Lid: 10.1601/ex.3006. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC 29543. High-quality 16S rRNA sequence S000363191 (RDP), M88159 (Genbank). GOLD ID: Gc00158. GCAT ID: 000452_GCAT. Entrez genome id: 445. Sequenced strain: DSM 1740 is type material. Genome sequencing is completed. Number of genomes of this species sequenced 1 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3006

Family "Hydrogenimonaceae" N4Lid DOI: 10.1601/nm.3864

Genus *Hydrogenimonas*^{VP} Takai et al. 2004. N4Lid DOI: 10.1601/nm.3865

Hydrogenimonas thermophila^{VP} Takai et al. 2004. Source of type material recommended for DOE sponsored genome sequencing by the JGI: ATCC BAA-737. High-quality 16S rRNA sequence S000252415 (RDP), AB105048 (Genbank). N4Lid DOI: 10.1601/nm.3866

Order Nautiliales^{VP} Miroshnichenko et al 2004. N4Lid DOI: 10.1601/nm.3867

Family Nautiliaceae^{VP} Miroshnichenko et al 2004. N4Lid DOI: 10.1601/nm.3868

Genus Nautilia^{VP} Miroshnichenko et al. 2002. GOLD ID: Gi00397. GCAT ID: 001255_GCAT. Entrez genome id: 13240. Sequenced strain: Am-H is from a non-type strain. Genome sequencing is incomplete. Number of genomes of this species sequenced 1 (GOLD) 1 (NCBI). N4Lid DOI: 10.1601/nm.3869

Nautilia lithotrophica^{VP} Miroshnichenko et al. 2002. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 13520. High-quality 16S rRNA sequence S000083785 (RDP), AJ404370 (Genbank). N4Lid DOI: 10.1601/nm.3870

Genus Caminibacter^{VP} Alain et al. 2002. N4Lid DOI: 10.1601/nm.3871

Caminibacter hydrogeniphilus^{VP} Alain et al. 2002. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 14510. High-quality 16S rRNA sequence S000131276 (RDP), AJ309655 (Genbank). N4Lid DOI: 10.1601/nm.3872

Caminibacter mediatlanticus^{VP} Voordeckers et al. 2005. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 16658. High-quality 16S rRNA sequence S000505089 (RDP), AY691430 (Genbank). GOLD ID: Gi01407. GCAT ID: 000863_GCAT. Sequenced strain: DSM 16658 is type material. Genome sequencing is incomplete. Number of genomes of this species sequenced 1 (GOLD). N4Lid 10.1601/nm.9362

Caminibacter profundus^{VP} Miroshnichenko et al. 2004. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 15016. High-quality 16S rRNA sequence S000264512 (RDP), AJ535664 (Genbank). N4Lid DOI: 10.1601/nm.3873

Genus Lebetimonas^{VP} Takai et al. 2005. N4Lid 10.1601/nm.8666

Lebetimonas acidiphila^{VP} Takai et al. 2005. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 16356. High-quality 16S rRNA sequence S000469711 (RDP), AB167820 (Genbank). N4Lid 10.1601/nm.8667

Genus Nitratifactor^{VP} Nakagawa et al. 2005. N4Lid 10.1601/nm.9374

Nitratifactor salsuginis^{VP} Nakagawa et al. 2005. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 16511. High-quality 16S rRNA sequence S000496333 (RDP), AB175500 (Genbank). N4Lid 10.1601/nm.9373

Genus Nitratiruptor^{VP} Nakagawa et al. 2005. N4Lid 10.1601/nm.9372

Nitratiruptor tergarcus^{VP} Nakagawa et al. 2005. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 16512. High-quality 16S rRNA sequence S000496332 (RDP), AB175499 (Genbank). N4Lid 10.1601/nm.9371

Genus Thioreductor^{VP} Nakagawa et al. 2005. N4Lid 10.1601/nm.9347

Thioreductor micantisoli^{VP} Nakagawa et al. 2005. Source of type material recommended for DOE sponsored genome sequencing by the JGI: DSM 16661. High-quality 16S rRNA sequence S000496331 (RDP), AB175498 (Genbank). N4Lid 10.1601/nm.9346