

Anaerococcus sp., Strain MJR7738A (Deposited as *Anaerococcus hydrogenalis*, Strain MJR7738A)

Catalog No. HM-1292

Product Description:

Anaerococcus sp., strain MJR7738A is a vaginal isolate obtained in 2014 from a pregnant woman with bacterial vaginosis in St. Louis, Missouri, USA. **HM-1292 was deposited to BEI Resources as *Anaerococcus hydrogenalis*; however, digital DNA-DNA hybridization (dDDH) analysis performed at BEI Resources resulted in reclassification to *Anaerococcus* sp.** HM-1292 was produced by inoculation of the deposited material into Modified Chopped Meat broth and grown for 2 days at 37°C in an anaerobic atmosphere (< 5% O₂; Remel™ Pack-Anaero™). Broth inoculum was passaged once in Modified Chopped Meat broth for 2 days at 37°C in an anaerobic atmosphere to produce this lot. Quality control testing was completed under propagation conditions unless otherwise noted.

Note: Quality control of HMP material is only performed to demonstrate that the material distributed by BEI Resources is identical to the deposited material. It should not be considered a complete characterization of the deposited organism.

Lot: 70009360

Manufacturing Date: 27OCT2017

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TEST	SPECIFICATIONS	RESULTS
Phenotypic Analysis Cellular morphology Colony morphology Motility (wet mount)	Gram-positive cocci Report results Report results	Gram-negative cocci Circular, convex, entire, smooth and white (Figure 1) Non-motile
Genotypic Analysis Sequencing of 16S ribosomal RNA gene (~ 760 base pairs) Digital DNA-DNA hybridization (dDDH) ¹	≥ 99% sequence identity to <i>A. hydrogenalis</i> , strain MJR7738 (GenBank: LRPL01000137.1) ≥ 70% for species identification	100% sequence identity to <i>A. hydrogenalis</i> , strain MJR7738 (GenBank: LRPL01000137.1) Not determined ^{2,3}
Purity (post-freeze) Anaerobic 7 days at 37°C on Tryptic Soy agar with 5% defibrinated sheep blood Aerobic with 5% CO ₂ 7 days at 37°C on Tryptic Soy agar with 5% defibrinated sheep blood	Growth consistent with expected colony morphology Growth consistent with expected colony morphology or no growth	Growth consistent with expected colony morphology No growth

¹Relatedness between bacterial strains has traditionally been determined using DDH. For additional information, refer to Auch, A. F., et al. "Digital DNA-DNA Hybridization for Microbial Species Delineation by Means of Genome-to-Genome Sequence Comparison." *Stand. Genomic Sci.* 2 (2010): 117-134. PubMed: 21304684.

²The whole genome of *Anaerococcus* sp., strain MJR7738 (contig total length approximately 2.1 megabase pairs) was sequenced using the Illumina® MiSeq® system and was analyzed and assembled with Kraken version 2.1.3 and Unicycler v0.5.1.

³dDDH testing rules out all species listed in Figure 2; however, this does not rule out species for which the type strains' whole genome sequences are not available. HM-1292 represents a potential new species.

Figure 1: Colony Morphology

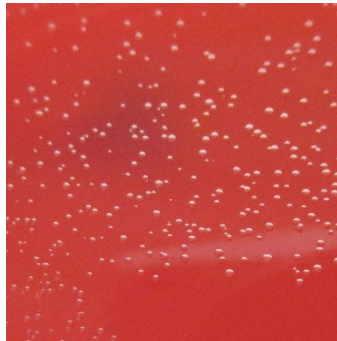


Figure 2: Digital DNA-DNA hybridization (dDDH)

Species	Strain	Accession Number	GGD ¹ vs. HM-1292 (Deposited as <i>Anaerococcus hydrogenalis</i>)
<i>Anaerococcus jeddahensis</i>	SB3	GCA_001182725	64.1%
<i>Peptoniphilus harei</i>	NCTC 13076	GCA_900454685	51.5%
<i>Anaerococcus rubeifantis</i>	mt16	GCA_001487145	50.4%
<i>Anaerococcus ihuae</i>	Marseille-Q5893 ^T	GCA_928368155	46.7%
<i>Anaerococcus lactolyticus</i>	ATCC [®] 51172 TM	GCA_000156575	37.2%
<i>Anaerococcus obesiensis</i>	ph10	GCA_000311745	35.5%
<i>Anaerococcus vaginalis</i>	ATCC [®] 51170 TM	GCA_000163295	34.9%
<i>Anaerococcus degeneri</i>	DSM 29674	GCA_017874475.1	34.5%
<i>Anaerococcus pacaensis</i>	9403502	GCA_000308255	31.5%
<i>Anaerococcus senegalensis</i>	JC48	GCA_000321005	31.0%
<i>Anaerococcus hydrogenalis</i>	DSM 7454	GCA_000173355	30.8%
<i>Anaerococcus porci</i>	DSM 101005	GCF_009696575	22.1%
<i>Anaerococcus faecalis</i>	AGMB00486	GCA_013371425	21.4%

¹Genome-to-genome distance (GGD)

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