

***Mycobacterium mantenii*, Strain NLA000401474T****Catalog No. NR-49079**

**Product Description:** *Mycobacterium mantenii* (*M. mantenii*), strain NLA000401474T was isolated in 2004 from a lymph node biopsy specimen from a 2-year-old female patient in the Netherlands.

**Lot<sup>1</sup>:** 64362428**Manufacturing Date:** 01AUG2016

TEST	SPECIFICATIONS	RESULTS
<b>Phenotypic Analysis<sup>2,3</sup></b>		
Cellular morphology	Rods	Rods
Colony morphology <sup>4</sup>	Report results	Circular, raised, entire, smooth and yellow (Figure 1)
Growth rate	≥ 7 days	11 days
Growth at 45°C	Negative	Negative
Growth at 55°C	Report results	Negative
Acid-fast stain	Positive (red colonies)	Positive (red colonies)
Pigmentation in the dark (Scotochromogen)	Positive	Positive
Photoinduction for 1 hour (Photochromogen)	Negative	Negative
Nonchromogen (no pigment)	Negative	Negative
Biochemical tests		
Catalase	Report results	Positive
Catalase (semiquantitative)	Positive	Positive
Catalase (68°C)	Positive	Positive
Iron uptake	Report results	Negative
Nitrate reduction	Negative	Negative
Tween 80 hydrolysis	Negative	Negative
Urease	Positive	Positive
Growth in the presence of 5% sodium chloride	Report results	Negative
Growth in the presence of thiophene-2-carboxylic acid hydrazide (TCH)	Positive	Positive
<b>Genotypic Analysis</b>		
Sequencing of 16S ribosomal RNA gene (~1470 base pairs)	≥ 99% sequence identity to <i>M. mantenii</i> type strain (GenBank: FJ042897.1)	99.9% sequence identity to <i>M. mantenii</i> type strain (GenBank: FJ042897.1)
Digital DNA-DNA hybridization (dDDH) <sup>5</sup>	≥ 70% for species identification	Not determined <sup>6,7</sup> (Table 1)
<b>Purity (post-freeze)</b>		
Middlebrook 7H10 agar with OADC enrichment <sup>8</sup>	Growth consistent with expected colony morphology	Growth consistent with expected colony morphology
Tryptic Soy agar <sup>8</sup>	Report results	Growth consistent with expected colony morphology
<b>Viability (post-freeze)<sup>4</sup></b>	Growth	Growth

<sup>1</sup>NR-49079 was produced by inoculation of the deposited material into Middlebrook 7H9 broth with ADC enrichment and grown for 14 days at 37°C in an aerobic atmosphere with 5% CO<sub>2</sub>. Broth inoculum was added to Middlebrook 7H10 agar with OADC enrichment kolles, which were grown for 6 days at 37°C in an aerobic atmosphere with 5% CO<sub>2</sub> to produce this lot.

<sup>2</sup>Information on Mycobacterium testing is available from Ribón, W. "Biochemical Isolation and Identification of Mycobacteria." *Biochemical Testing*, (2012) Jose C. Jimenez-Lopez (Ed.), InTech, <http://www.intechopen.com/books/biochemical-testing/biochemical-isolation-and-identification-of-mycobacteria> and Lévy-Frébault, V. V. and F. Portaels. "Proposed Minimal Standards for the Genus *Mycobacterium* and for Description of New Slowly Growing *Mycobacterium* Species." *Int. J. Syst. Bacteriol.* 42 (1992): 315-323. PubMed: 1581193.

<sup>3</sup>Phenotypic characterization of *M. mantenii* was performed following van Ingen, J., et al. "*Mycobacterium mantenii* sp. nov., a Pathogenic, Slowly Growing, Scotochromogenic Species." *Int. J. Syst. Evol. Microbiol.* 59 (2009): 2782-2787. PubMed: 19625425.

<sup>4</sup>11 days at 37°C in an aerobic atmosphere with 5% CO<sub>2</sub> on Middlebrook 7H10 agar with OADC enrichment

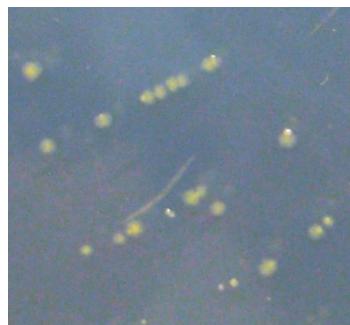
<sup>5</sup>Relatedness between bacterial strains has traditionally been determined using DDH. For additional information, refer to Auch, A.F., et al. "Digital DNA-DNA Hybridization for Microbial Species Delineation by Means of Genome-to-Genome Sequence Comparison." *Stand. Genomic Sci.* 2 (2010): 117-134. PubMed: 21304684.

<sup>6</sup>The whole genome of *M. mantenii*, strain NLA000401474T (Contig Total Length ~ 6.1 megabase pairs) was sequenced using the Illumina® MiSeq® system and was assembled and analyzed with CLC Genomics Workbench Version 7.0.2.

<sup>7</sup>The required whole genome sequence for the type strain of this species is not available. dDDH testing rules out all species listed in Table 1, however, this does not rule out species for which the type strains whole genome sequences are not available.

<sup>8</sup>Purity of this lot was assessed for 11 days at 37°C in an aerobic atmosphere with 5% CO<sub>2</sub>.

**Figure 1: Colony Morphology**



**Table 1: Digital DNA-DNA hybridization (dDDH)**

Species	Strain	Accession #	GGD vs. NR-49079 (Deposited as: <i>M. mantenii</i> )
<i>M. abscessus</i> subsp. <i>abscessus</i>	Hauduroy L948 <sup>T</sup>	NC_010397.1	19.6
<i>M. abscessus</i> subsp. <i>bolletii</i>	BD <sup>T</sup>	AHAS00000000.1	19.5
<i>M. abscessus</i> subsp. <i>massiliense</i>	CCUG 48898 <sup>T</sup>	NZ_AP014547.1	19.4
<i>M. aromaticivorans</i>	JS19b1 <sup>T</sup>	JALN00000000.2	20.1
<i>M. aurum</i>	ATCC 23366 <sup>T</sup>	CVQQ01000001.1	20.1
<i>M. austroafricanum</i>	E9789-SA12441 <sup>T</sup>	HG964450.1	19.9
<i>M. avium</i> subsp. <i>avium</i>	ATCC 25291 <sup>T</sup>	ACFI00000000.1	29.8
<i>M. avium</i> subsp. <i>paratuberculosis</i>	ATCC 19698 <sup>T</sup>	AGAR00000000.1	30.3
<i>M. avium</i> subsp. <i>silvaticum</i>	6409 <sup>T</sup>	AYOC00000000.1	30.2
<i>M. bohemicum</i>	CIP 105808 <sup>T</sup>	CSTD01000001.1	24.9
<i>M. canariensis</i>	502329 <sup>T</sup>	BCSY00000000.1	20.2
<i>M. celatum</i>	ATCC 51131 <sup>T</sup>	BBUN00000000.1	22.4
<i>M. chelonae</i>	CM 6388 <sup>T</sup>	CP010946.1	19
<i>M. chlorophenicolum</i>	PCP-I <sup>T</sup>	JYNL00000000.1	20
<i>M. chubuense</i>	48013 <sup>T</sup>	NC_018027.1	19.9
<i>M. colombiense</i>	10B <sup>T</sup>	AFVW00000000.2	36.6
<i>M. conceptionense</i>	D16 <sup>T</sup>	CTEF00000000.1	20.2
<i>M. cosmeticum</i>	LTA-388 <sup>T</sup>	CCBB00000000.1	20.4
<i>M. crocinum</i>	czh-42 <sup>T</sup>	BBHD00000000.1	21.7
<i>M. farcinogenes</i>	IEMVT 75 <sup>T</sup>	CCAY00000000.1	20
<i>M. fluoranthrenivorans</i>	FA4 <sup>T</sup>	BBFT00000000.1	21.6
<i>M. fortuitum</i> subsp. <i>fortuitum</i>	ATCC 6841 <sup>T</sup>	CP014258.1	19.7
<i>M. fortuitum</i> subsp. <i>acetamidolyticum</i>	NCH E11620 <sup>T</sup>	BCSZ00000000.1	19.8
<i>M. gastri</i>	ATCC 15754 <sup>T</sup>	AZYN00000000.1	22.5
<i>M. genavense</i>	2289 <sup>T</sup>	JAGZ00000000.1	23.5
<i>M. haemophilum</i>	ATCC 29548 <sup>T</sup>	CP011883.2	22.4
<i>M. hassiacum</i>	3849 <sup>T</sup>	ARBU00000000.1	20.3
<i>M. hodleri</i>	EMI2 <sup>T</sup>	BBGO00000000.1	23.1
<i>M. intracellularare</i>	ATCC 13950 <sup>T</sup>	NC_016946.1	31.2
<i>M. kansasii</i>	ATCC 12478 <sup>T</sup>	NC_022663.1	22.3
<i>M. kyorinense</i>	KUM 060204 <sup>T</sup>	BBKA00000000.1	21.9

## Certificate of Analysis for NR-49079

Species	Strain	Accession #	GGD vs. NR-49079 (Deposited as: <i>M. mantenii</i> )
<i>M. mageritense</i>	938 <sup>T</sup>	CCBF00000000.1	20.1
<i>M. neoaurum</i>	ATCC 25795 <sup>T</sup>	JMDW00000000.1	19.9
<i>M. neworleansense</i>	W6705 <sup>T</sup>	CWKH00000000.1	20
<i>M. novocastrense</i>	73 <sup>T</sup>	BCTA00000000.1	20.2
<i>M. obuense</i>	47001 <sup>T</sup>	JYNU00000000.1	20.2
<i>M. pallens</i>	czh-8 <sup>T</sup>	BBHE00000000.1	21.9
<i>M. parascrofulaceum</i>	HSC-68 <sup>T</sup>	ADNV00000000.1	26.3
<i>M. pseudoshottsii</i>	L15 <sup>T</sup>	BCND00000000.1	21.3
<i>M. pyrenivorans</i>	17A3 <sup>T</sup>	BBHB00000000.1	22.1
<i>M. rufum</i>	JS14 <sup>T</sup>	JROA00000000.1	20
<i>M. rutilum</i>	czh-117 <sup>T</sup>	BBHF00000000.1	23.6
<i>M. septicum</i>	W4964 <sup>T</sup>	CBMO000000000.1	19.9
<i>M. setense</i>	ABO-M06 <sup>T</sup>	JTJW00000000.1	19.9
<i>M. simiae</i>	ATCC 25275 <sup>T</sup>	CBMJ000000000.2	22.9
<i>M. smegmatis</i>	ATCC 19420 <sup>T</sup>	LN831039.1	19.9
<i>M. thermoresistibile</i>	ATCC 19527 <sup>T</sup>	BCTB00000000.1	20
<i>M. triplex</i>	90-1019 <sup>T</sup>	CCAU000000000.1	24.2
<i>M. tuberculosis</i>	H37Rv <sup>T</sup>	NC_000962.3	22
<i>M. vaccae</i>	ATCC 15483 <sup>T</sup>	BCRS00000000.1	20.3
<i>M. vanbaalenii</i>	PYR-1 <sup>T</sup>	NC_008726.1	20.1
<i>M. vulneris</i>	NLA000700772 <sup>T</sup>	CCBG000000000.1	20.1
<i>M. yongonense</i>	05-1390 <sup>T</sup>	NC_021715.1	30.9
<i>Nocardia asteroides</i>	NBRC 15531 <sup>T</sup>	BAFO00000000.2	19.3

Date: 15 JUN 2017

Signature:



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