

Certificate of Analysis for NR-36439

Paenibacillus barengoltzii, Strain G22

Catalog No. NR-36439

Product Description:

Paenibacillus barengoltzii (P. barengoltzii), strain G22 was isolated from mouse intestine in the United States.

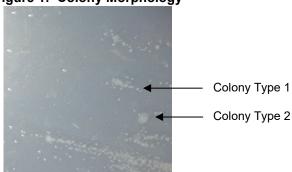
Lot: 70012206¹ Manufacturing Date: 05FEB2018

TEST	SPECIFICATIONS	RESULTS
Phenotypic Analysis		
Cellular morphology	Gram-positive rods	Gram-positive rods
Colony morphologies ^{2,3}	Report results	Colony type 1: Circular, raised, entire, smooth and cream (Figure 1)
		Colony type 2: Circular, low convex, entire, opaque and cream (Figure 1)
Motility	Report results	Motile
VITEK® MS (MALDI-TOF)4	Paenibacillus spp.	Paenibacillus spp. (≥ 84%)
Genotypic Analysis		
Sequencing of 16S ribosomal RNA gene (~ 800 base pairs)	≥ 99% sequence identity to <i>P. barengoltzii</i> , strain G22 (GenBank: ASSZ01000038.1)	100% sequence identity to P. barengoltzii, strain G22 (GenBank: ASSZ01000038.1)
Digital DNA-DNA hybridization (dDDH) ⁵	≥ 70% for species identification ′	P. barengoltzii (79.4%)
Purity (post-freeze) ⁶	Consistent with expected colony morphology	Consistent with expected colony morphology
Viability (post-freeze) ²	Growth	Growth

¹NR-36439 was produced by inoculation of the deposited material into Nutrient broth and grown 3 days at 30°C in an aerobic atmosphere. Broth inoculum was added to Nutrient agar kolles which were grown 4 days at 30°C in an aerobic atmosphere to produce this lot.

⁶Purity of this lot was assessed for 7 days at 37°C in an aerobic atmosphere with 5% CO₂ on Nutrient agar.





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²2 days at 30°C in an aerobic atmosphere on Nutrient agar

³Two colony types were observed. Plating of the individual colony types showed that colony type 1 reverted to the mixed colony type and colony type 2 did not revert.

⁴VITEK[®] MS (MALDI-TOF) was used to confirm to genus.

⁵Relatedness 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." <u>Stand. Genomic Sci.</u> 2 (2010): 117-134. PubMed: 21304684. *P. barengoltzii*, strain NBRC 101215^T (GenBank: BILV00000000.1) was used for dDDH analysis.



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/Heather Couch/ Heather Couch

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Program Manager or designee, ATCC Federal Solutions

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