

***Pseudomonas aeruginosa*, Strain MRSN 16344**

Catalog No. NR-51581

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Product Description:

Pseudomonas aeruginosa (*P. aeruginosa*), strain MRSN 16344 was isolated in 2013 from a human wound as part of a surveillance program in the United States. *P. aeruginosa*, strain MRSN 16344 was deposited as sensitive to aztreonam, ceftazidime and piperacillin/tazobactam, intermediately resistant to amikacin, cefepime and meropenem and resistant to ciprofloxacin, gentamicin, imipenem, levofloxacin and tobramycin.

Lot: 70025068¹

Manufacturing Date: 03JUL2019

TEST	SPECIFICATIONS	RESULTS
Phenotypic Analysis Cellular morphology Colony morphology ² Motility (wet mount) VITEK [®] 2 (GN card)	Gram-negative rods Report results Report results <i>P. aeruginosa</i> (≥ 89%)	Gram-negative rods Punctiform (Figure 1) Motile <i>P. aeruginosa</i> (≥ 97%)
Antibiotic Susceptibility Profile³ VITEK [®] (AST-GN81 Card) Ampicillin Amoxicillin/clavulanic acid Piperacillin/tazobactam Cefazolin Cefoxitin Ceftazidime Ceftriaxone Cefepime Meropenem Amikacin Gentamicin Tobramycin Ciprofloxacin Levofloxacin Tetracycline Nitrofurantoin Trimethoprim/sulfamethoxazole	Report results Report results Sensitive Report results Report results Sensitive Report results Intermediate Intermediate Intermediate Resistant Resistant Resistant Resistant Report results Report results Report results	Resistant (≥ 32 µg/mL) Resistant (≥ 32 µg/mL) Sensitive (≤ 8 µg/mL) Resistant (≥ 64 µg/mL) Resistant (≥ 32 µg/mL) Sensitive (≤ 1 µg/mL) Resistant (≥ 64 µg/mL) Sensitive (≤ 8 µg/mL) ⁴ Sensitive (1 µg/mL) ⁵ Sensitive (16 µg/mL) ⁴ Intermediate (8 µg/mL) ⁴ Sensitive (≤ 4 µg/mL) ⁶ Intermediate (2 µg/mL) ⁷ Intermediate (4 µg/mL) ⁸ Resistant (≥ 16 µg/mL) Resistant (≥ 512 µg/mL) ≥ 320 µg/mL ⁹
Genotypic Analysis Sequencing of 16S ribosomal RNA gene (~ 1420 base pairs)	≥ 99% sequence identity to <i>P. aeruginosa</i> , strain MRSN 16344 (GenBank: RXVS01000152.1)	99.9% sequence identity to <i>P. aeruginosa</i> , strain MRSN 16344 (GenBank: RXVS01000152.1)
Purity (post-freeze)¹⁰	Growth consistent with expected colony morphology	Growth consistent with expected colony morphology
Viability (post-freeze)²	Growth	Growth

¹NR-51581 was produced by inoculation of the depositor material into Tryptic Soy broth and grown for 1 day at 37°C in an aerobic atmosphere. Broth inoculum was added to Tryptic Soy agar kolles, which were grown for 1 day at 37°C in an aerobic atmosphere to produce this lot.

²1 day at 37°C in an aerobic atmosphere on Tryptic Soy agar

³Minimum Inhibitory Concentration (MIC); MIC Interpretation Guideline: CLSI M100-S28 (2018)

⁴Susceptibility results for this antibiotic are within one doubling dilution of specification, which is considered an equivalent result.

⁵*P. aeruginosa*, strain MRSN 16344 was deposited as intermediately resistant to meropenem. Repeated antibiotic susceptibility testing determined that strain MRSN 16344 is sensitive to meropenem.

⁶*P. aeruginosa*, strain MRSN 16344 was deposited as resistant to tobramycin. Repeated antibiotic susceptibility testing determined that strain MRSN 16344 is sensitive to tobramycin.

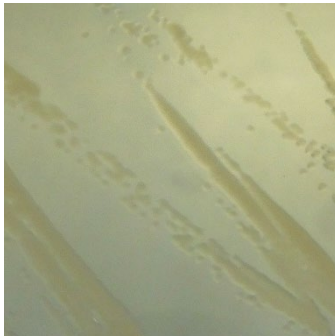
⁷*P. aeruginosa*, strain MRSN 16344 was deposited as resistant to ciprofloxacin. Repeated antibiotic susceptibility testing determined that strain MRSN 16344 is intermediately resistant to ciprofloxacin.

⁸*P. aeruginosa*, strain MRSN 16344 was deposited as resistant to levofloxacin. Repeated antibiotic susceptibility testing determined that strain MRSN 16344 is intermediately resistant to levofloxacin.

⁹Trimethoprim/sulfamethoxazole MIC interpretive standards are not available for *P. aeruginosa*, however most clinical isolates are resistant to trimethoprim/sulfamethoxazole. For more information, please refer to Köhler, T., et al. "Multidrug Efflux in Intrinsic Resistance to Trimethoprim and Sulfamethoxazole in *Pseudomonas aeruginosa*." *Antimicrob. Agents Chemother.* 40 (1996): 2288-2290. PubMed: 9036831.

¹⁰Purity of this lot was assessed for 8 days at 37°C in an aerobic atmosphere with and without 5% CO₂ on Tryptic Soy agar.

Figure 1: Colony Morphology



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11 DEC 2019

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