

***Pseudomonas aeruginosa*, Strain MRSN 6695**

Catalog No. NR-51552

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Product Description:

Pseudomonas aeruginosa (*P. aeruginosa*), strain MRSN 6695 was isolated in 2012 from human urine as part of a surveillance program in the United States. *P. aeruginosa*, strain MRSN 6695 was deposited as sensitive to amikacin, gentamicin and tobramycin and resistant to aztreonam, ceftazidime, ciprofloxacin, imipenem, levofloxacin, meropenem and piperacillin/tazobactam with intermediate resistance to cefepime.

Lot: 70024979¹

Manufacturing Date: 09MAY2019

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 Circular, low convex, entire, opaque and cream (Figure 1) Motile <i>P. aeruginosa</i> (99%)
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 Resistant Report results Report results Resistant Report results Intermediate Resistant Sensitive Sensitive Sensitive Resistant Resistant Report results Report results Report results	Resistant (≥ 32 µg/mL) Resistant (≥ 32 µg/mL) Resistant (≥ 128 µg/mL) Resistant (≥ 64 µg/mL) Resistant (≥ 64 µg/mL) Intermediate (16 µg/mL) ⁴ Resistant (≥ 64 µg/mL) Sensitive (8 µg/mL) ⁵ Intermediate (4 µg/mL) ⁶ Sensitive (8 µg/mL) Sensitive (2 µg/mL) Sensitive (≤ 1 µg/mL) Sensitive (1 µg/mL) ⁷ Intermediate (4 µg/mL) ⁸ Resistant (≥ 16 µg/mL) Resistant (≥ 512 µg/mL) 80 µg/mL ⁹
Genotypic Analysis Sequencing of 16S ribosomal RNA gene (~ 1470 base pairs)	≥ 99% sequence identity to <i>P. aeruginosa</i> , strain MRSN 6695 (GenBank: RXTJ01000040.1)	99.9% sequence identity to <i>P. aeruginosa</i> , strain MRSN 6695 (GenBank: RXTJ01000040.1)
Purity (post-freeze)¹⁰	Growth consistent with expected colony morphology	Growth consistent with expected colony morphology
Viability (post-freeze)²	Growth	Growth

¹NR-51552 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)

⁴*P. aeruginosa* strain MRSN 6695 was deposited as resistant to ceftazidime. Repeated antibiotic susceptibility testing determined that strain MRSN 6695 is intermediately resistant to ceftazidime.

⁵The susceptibility result for this antibiotic is within one doubling dilution of specification, which is considered an equivalent result.

⁶*P. aeruginosa* strain MRSN 6695 was deposited as resistant to meropenem. Repeated antibiotic susceptibility testing determined that strain MRSN 6695 is intermediately resistant to meropenem.

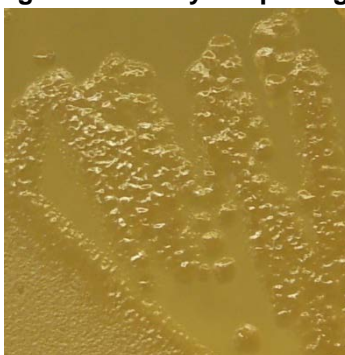
⁷*P. aeruginosa* strain MRSN 6695 was deposited as resistant to ciprofloxacin. Repeated antibiotic susceptibility testing determined that strain MRSN 6695 is sensitive to ciprofloxacin.

⁸*P. aeruginosa* strain MRSN 6695 was deposited as resistant to levofloxacin. Repeated antibiotic susceptibility testing determined that strain MRSN 6695 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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07 JAN 2020

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