

Pseudomonas aeruginosa, Strain MRSN 25678

Catalog No. NR-51599

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Product Description:

Pseudomonas aeruginosa (*P. aeruginosa*), strain MRSN 25678 was isolated from human urine in 2014 as part of a surveillance program in the United States. *P. aeruginosa*, strain MRSN 25678 was deposited as sensitive to amikacin, imipenem and tobramycin and resistant to aztreonam, ciprofloxacin, cefepime and levofloxacin, with intermediate resistance to ceftazidime, gentamicin, meropenem and piperacillin/tazobactam.

Lot: 70025104¹

Manufacturing Date: 01AUG2020

| 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, convex, entire, smooth and cream (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 Etest® antibiotic test strips ⁹ Gentamicin | Report results Report results Intermediate Report results Report results Intermediate Report results Resistant Intermediate Sensitive Intermediate Sensitive Resistant Resistant Report results Report results Report results Intermediate | Resistant (≥ 32 µg/mL) Resistant (≥ 32 µg/mL) Sensitive (16 µg/mL) ⁴ Resistant (≥ 64 µg/mL) Resistant (≥ 64 µg/mL) Intermediate (16 µg/mL) Resistant (≥ 64 µg/mL) Resistant (≥ 64 µg/mL) Intermediate (2-4 µg/mL) Intermediate (32 µg/mL) ⁵ Sensitive (4 µg/mL) ⁴ Sensitive (≤ 1 µg/mL) Intermediate (2 µg/mL) ⁶ Intermediate (4 µg/mL) ⁷ Resistant (≥ 16 µg/mL) Resistant (≥ 512 µg/mL) ≥ 320 µg/mL ⁸ Intermediate (12 µg/mL) |
| Genotypic Analysis Sequencing of 16S ribosomal RNA gene (~ 1400 base pairs) | ≥ 99% sequence identity to <i>P. aeruginosa</i> , strain MRSN 25678 (GenBank: RXUN01000193.1) | 100% sequence identity to <i>P. aeruginosa</i> , strain MRSN 25678 (GenBank: RXUN01000193.1) |
| Purity (post-freeze)¹⁰ | Growth consistent with expected colony morphology | Growth consistent with expected colony morphology |
| Viability (post-freeze)² | Growth | Growth |

¹NR-51599 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)

⁴The susceptibility result for this antibiotic is within one doubling dilution of specification, which is considered an equivalent result.

⁵*P. aeruginosa*, strain MRSN 25678 was deposited as sensitive to amikacin, but showed a MIC of 32 µg/mL (interpreted as intermediate) for amikacin during QC testing. Testing was performed in duplicate.

⁶*P. aeruginosa*, strain MRSN 25678 was deposited as resistant to ciprofloxacin, but showed a MIC of 2 µg/mL (interpreted as intermediate) for ciprofloxacin during QC testing. Testing was performed in duplicate.

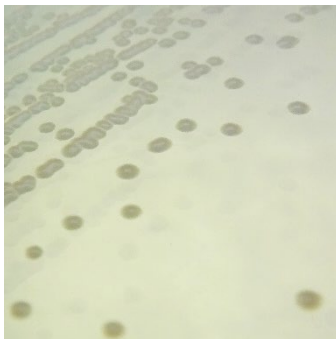
⁷*P. aeruginosa*, strain MRSN 25678 was deposited as resistant to levofloxacin, but showed a MIC of 4 µg/mL (interpreted as intermediate) for levofloxacin during QC testing. Testing was performed in duplicate.

⁸Trimethoprim/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.

⁹1 day at 37°C in an aerobic atmosphere on Mueller Hinton agar

¹⁰Purity of this lot was assessed for 7 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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Program Manager or designee, ATCC Federal Solutions

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