

***Klebsiella pneumoniae*, Strain MRSN 368001**

Catalog No. NR-55546

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Contributor:

Multidrug-Resistant Organism Repository and Surveillance Network (MRSN), Bacterial Disease Branch, Walter Reed Army Institute of Research, Silver Spring, Maryland, USA

Manufacturer:

BEI Resources

Product Description:

Bacteria Classification: *Enterobacteriaceae*, *Klebsiella*

Species: *Klebsiella pneumoniae*

Strain: MRSN 368001

Original Source: *Klebsiella pneumoniae* (*K. pneumoniae*), strain MRSN 368001 was isolated in 2015 from a human wound sample in Africa as part of a global surveillance program.¹

Comments: *K. pneumoniae*, strain MRSN 368001 was deposited as part of the MRSN *Klebsiella pneumoniae* Diversity Panel available from BEI Resources as NR-55604. NR-55546 was deposited as multi-locus sequence type (MLST) ST 14, K-locus type (KL) 2, O-locus type (OL) O1v1, VIR score 0. In addition, strain MRSN 368001 has a glycine-aspartic acid (GD) insertion in the B-strand loop of the OmpK36 protein (OmpK36GD). MRSN 368001 was deposited as an extensively drug-resistant strain (XDR), sensitive to tetracycline and tigecycline and resistant to amikacin, ampicillin/sulbactam, aztreonam, cefepime, ceftazidime, ceftriaxone, ceftazidime/avibactam, ceftolozane/tazobactam, ciprofloxacin, ertapenem, meropenem, gentamicin, imipenem, levofloxacin, piperacillin/tazobactam, tobramycin and trimethoprim/sulfamethoxazole. Strain MRSN 368001 is reported to have three aminoglycoside 3'-phosphotransferase genes [*aac*(6')-Ib-cr5, *aadA2* and *aph*(3')-VI; conferring resistance to various aminoglycosides], one 16S rRNA methyltransferase gene (*armA*; conferring resistance to aminoglycoside antibiotics), four beta-lactamase genes (*bla*_{CTX-M-15}, *bla*_{SHV-28}, *bla*_{NDM-1} and *bla*_{OXA-1}; conferring resistance to beta-lactams), two chloramphenicol acetyltransferase genes (*catA1* and *catB3*; conferring resistance to chloramphenicol), one fosfomycin resistance gene (*fosA_{gen}*; conferring resistance to fosfomycin), one macrolide phosphotransferase gene [*mph*(E); conferring resistance to macrolides], one quinolone resistance gene (*qnrB1*; conferring resistance to quinolones), one ABC-F subfamily protein gene [*msr*(E); conferring resistance to erythromycin and streptogramin B antibiotics], one sulfonamide resistance gene (*su1*; conferring resistance to sulfonamides) and two dihydrofolate reductase genes (*dhfrA1* and *dhfrA12*; conferring

resistance to trimethoprim).¹ The complete genome of *K. pneumoniae*, strain MRSN 368001 has been sequenced (GenBank: [JAGYDN000000000](https://www.ncbi.nlm.nih.gov/nuccore/JAGYDN000000000)).

K. pneumoniae is a Gram-negative enterobacterium that is a major cause of nosocomial infections of the urinary and respiratory tracts. Due to the extensive spread of antibiotic-resistant strains, especially of extended-spectrum β -lactamase (ESBL)-producing strains, there has been renewed interest in *Klebsiella* infections.^{2,3,4}

Material Provided:

Each vial contains approximately 0.5 mL of bacterial culture in Tryptic Soy broth supplemented with 10% glycerol.

Note: If homogeneity is required for your intended use, please purify prior to initiating work.

Packaging/Storage:

NR-55546 was packaged aseptically in cryovials. The product is provided frozen and should be stored at -60°C or colder immediately upon arrival. For long-term storage, the vapor phase of a liquid nitrogen freezer is recommended. Freeze-thaw cycles should be avoided.

Growth Conditions:

Media:

Nutrient broth or Tryptic Soy broth or equivalent
Nutrient agar or Tryptic Soy agar or Tryptic Soy agar with 5% defibrinated sheep blood or equivalent

Incubation:

Temperature: 37°C
Atmosphere: Aerobic

Propagation:

1. Keep vial frozen until ready for use, then thaw.
2. Transfer the entire thawed aliquot into a single tube of broth.
3. Use several drops of the suspension to inoculate an agar slant and/or plate.
4. Incubate the tube, slant and/or plate at 37°C for 1 day.

Citation:

Acknowledgment for publications should read "The following reagent was obtained through BEI Resources, NIAID, NIH: *Klebsiella pneumoniae*, Strain MRSN 368001, NR-55546. This strain is part of the *Klebsiella pneumoniae* Diversity Panel provided by the Multidrug-Resistant Organism Repository and Surveillance Network (MRSN) at the Walter Reed Army Institute of Research (WRAIR)."

Biosafety Level: 2

Appropriate safety procedures should always be used with this material. Laboratory safety is discussed in the following publication: U.S. Department of Health and Human Services, Public Health Service, Centers for Disease Control and Prevention, and National Institutes of Health. [Biosafety in Microbiological and Biomedical Laboratories](#). 6th ed. Washington, DC: U.S. Government Printing Office, 2020; see www.cdc.gov/biosafety/publications/bmb15/index.htm.

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References:

1. McGann, P., Personal Communication.
2. Lascols, C., et al. "Increasing Prevalence and Dissemination of NDM-1 Metallo-β-Lactamase in India: Data from the SMART Study (2009)." J. Antimicrob. Chemother. 66 (2011): 1992-1997. PubMed: 21676902.
3. Ramirez, M. S., et al. "Multidrug-Resistant (MDR) *Klebsiella pneumoniae* Clinical Isolates: A Zone of High Heterogeneity (HHZ) as a Tool for Epidemiological Studies." Clin. Microbiol. Infect. 18 (2012): E254-E258. PubMed: 22551038.
4. Podschun, R. and U. Ullmann. "*Klebsiella* spp. as Nosocomial Pathogens: Epidemiology, Taxonomy, Typing Methods, and Pathogenicity Factors." Clin. Microbiol. Rev. 11 (1998): 589-603. PubMed: 9767057.

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