

# **Product Information Sheet for NR-9318**

# Genomic DNA from Burkholderia mallei, Strain China 7 (NBL 7)

## Catalog No. NR-9318

### For research use only. Not for human use.

#### Contributor:

Herbert P. Schweizer, Ph.D., Department of Microbiology, Immunology and Pathology, College of Veterinary Medicine and Biomedical Sciences, Colorado State University, Fort Collins, Colorado

### **Product Description:**

Genomic DNA was isolated from a preparation of Burkholderia mallei (B. mallei), strain China 7 (BEI Resources NR-4071). NR-4071 was derived from ATCC 23344™ via several passages by different individuals prior to its deposit at BEI Resources. Genome variability upon passage has been reported to be a feature of B. mallei, strain China 7 (ATCC<sup>®</sup> 23344<sup>TM</sup>). Genomic DNA from BEI Resources NR-23 (a preparation of *B. mallei* strain China 7 that was produced directly from ATCC<sup>®</sup> 23344<sup>TM</sup>) is available as BEI Resources NR-2535.

B. mallei, strain China 7 was isolated from postmortem cultures of knee fluid, skin pustules, and blood of a Chinese soldier who died in Burma (1944) from a glanders-melioidosis type of infection. The complete genomic sequence of Burkholderia mallei, strain China 7 has been determined (GenBank: CP000010 and CP000011).2

NR-9318 has been qualified for PCR applications by amplification of approximately 1500 bp of the 16S ribosomal RNA gene.

### **Material Provided:**

Each vial contains approximately 4 to 6 µg of bacterial genomic DNA in TE buffer (10 mM Tris-HCl, 1 mM EDTA, pH ~ 7.4). The concentration is shown on the Certificate of Analysis. The vial should be centrifuged prior to opening.

#### Packaging/Storage:

NR-9318 was packaged aseptically in screw-capped plastic cryovials. The product is provided frozen and should be stored at 4°C or colder immediately upon arrival. For optimal long-term storage, freezing the material at -20°C or colder is recommended. Freeze-thaw cycles should be minimized.

#### Citation:

Acknowledgment for publications should read "The following reagent was obtained through the NIH Biodefense and Emerging Infections Research Resources Repository, NIAID. NIH: Genomic DNA from Burkholderia mallei, Strain China 7 (NBL 7), NR-9318."

### Biosafety Level: 1

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. 5th ed. Washington, DC: U.S. Government Printing Office, 2007; see www.cdc.gov/od/ohs/biosfty/bmbl5/bmbl5toc.htm.

#### **Disclaimers:**

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#### References:

- 1. Romero, C. M., et al. "Genome Sequence Alterations Detected upon Passage of Burkholderia mallei ATCC 23344 in Culture and in Mammalian Hosts." BMC Genomics 7 (2006): 228-238. PubMed: 16953889.
- Nierman, W. C., et al. "Structural Flexibility in the Burkholderia mallei Genome." Proc. Natl. Acad. Sci. U.S.A. 101 (2004): 14246-14251. PubMed: 15377793. GenBank: CP000010 and CP000011.

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P.O. Box 4137 Manassas, VA 20108-4137 USA www.beiresources.org

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- Bauernfeind, A., et al. "Molecular Procedure for Rapid Detection of Burkholderia mallei and Burkholderia pseudomallei." J. Clin. Microbiol. 36 (1998): 2737-2741. PubMed: 9705426.
- Godoy, D., et al. "Multilocus Sequence Typing and Evolutionary Relationships Among the Causative Agents of Melioidosis and Glanders, *Burkholderia pseudomallei* and *Burkholderia mallei*." J. Clin. Microbiol. 41 (2003): 2068-2079. PubMed: 12734250.
- Gee, J. E., et al. "Use of 16S rRNA Gene Sequencing for Rapid Identification and Differentiation of *Burkholderia* pseudomallei and *B. mallei*." <u>J. Clin. Microbiol.</u> 41 (2003): 4647-4654. PubMed: 14532197.
- Ong, C., et al. "Patterns of Large-Scale Genomic Variation in Virulent and Avirulent *Burkholderia* Species." <u>Genome Res.</u> 14 (2004): 2295-2307. PubMed: 15520292.

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800-359-7370

Fax: 703-365-2898

E-mail: contact@beiresources.org