

***Borrelia burgdorferi*, Signature-Tagged Mutagenesis Library Clone T03TC121 (Gene IR\_BB\_L35-BB\_L36)**

**Catalog No. NR-22408**

**For research use only. Not for use in humans.**

**Contributor:**

Steven J. Norris, Ph.D., Professor and Vice Chair for Research, Department of Pathology and Laboratory Medicine, University of Texas Health Science Center at Houston Medical School, Houston, Texas, USA

**Manufacturer:**

BEI Resources

**Product Description:**

Bacteria Classification: Spirochaetaceae, *Borrelia*

Species: *Borrelia burgdorferi*

Strain: B31, clone 5A18NP1

Signature-Tagged Mutagenesis Library Clone: T03TC121

Replicon: Circular plasmid cp32-8

Gene: IR\_BB\_L35-BB\_L36 (intergenic region)

Insertion Site<sup>1,2</sup>: 22910

Original Source: *Borrelia burgdorferi* (*B. burgdorferi*), clone T03TC121 was produced by signature-tagged mutagenesis (STM) of the intergenic region between the BB\_L35 and BB\_L36 genes.<sup>1,2</sup>

Comments: *B. burgdorferi*, strain B31 5A18NP1 STM library clone T03TC121 lacks linear plasmids lp5, lp28-4 and lp56. The plasmid profile was determined by PCR using plasmid-specific primers.<sup>2</sup>

*B. burgdorferi* is a Gram-negative, motile spirochete.<sup>3</sup> It is a zoonotic, vector-borne pathogen transmitted by ticks and the etiologic agent of Lyme disease, now the most common tick-transmitted disease in the United States.<sup>4</sup> *B. burgdorferi* is predominant in North America, but also exists in Europe.

*B. burgdorferi*, strain B31 was originally isolated in 1981 from adult ticks (*Ixodes dammini*) collected from lower vegetation on Shelter Island, New York, USA.<sup>3,4</sup> Strain B31 is composed of a 910 kilobase (kb) linear chromosome, 9 circular plasmids (cp) and 12 linear plasmids (lp). Plasmids range in size from 5 kb to 56 kb and total 610 kb.<sup>2,5</sup> Continuous passage of *B. burgdorferi* is known to cause spontaneous loss of plasmids. The complete genome of *B. burgdorferi*, strain B31 has been sequenced (GenBank: [AE000783](https://www.ncbi.nlm.nih.gov/nuccore/AE000783)).<sup>6</sup>

*B. burgdorferi*, strain B31, clone 5A18NP1 was derived from the low-passage clone 5A18 of strain B31.<sup>7</sup> Clone 5A18NP1 lacks lp56 and lp28-4 and the BBE02 gene (a putative restriction-modification gene on lp25) was disrupted by homologous recombination resulting in kanamycin resistance.<sup>8</sup> Inactivation of BBE02 results in increased transformation efficiency and therefore, clone 5A18NP1, was used to create the STM library through the *mariner*-based transposition suicide *Himar1* delivery vector, pMarGent,

containing *aacC1* which confers gentamicin resistance.<sup>1,2,9</sup> STM is a negative selection method that identifies clones by unique DNA sequences that are incorporated into the transposable element.<sup>2</sup>

**Material Provided:**

Each vial contains approximately 0.5 mL of bacterial culture in Revised Barbour-Stoenner-Kelly medium supplemented with 200 micrograms per milliliter kanamycin, 40 micrograms per milliliter gentamicin and 15% glycerol.

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

**Packaging/Storage:**

NR-22408 was packaged aseptically in cryovials. The product is provided frozen and should be stored at -80°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:

Revised Barbour-Stoenner-Kelly broth (see Appendix I) with 200 micrograms per milliliter kanamycin and 40 micrograms per milliliter gentamicin

Revised Barbour-Stoenner-Kelly agar (see Appendix I) with 200 micrograms per milliliter kanamycin, 40 micrograms per milliliter gentamicin and 0.8% agar

Incubation:

Temperature: 32°C to 34°C (growth at 37°C may result in plasmid loss)<sup>1</sup>

Atmosphere: Microaerophilic (slower growth occurs under aerobic conditions<sup>1</sup>)

Propagation:

1. Keep vial in dry ice during inoculations.
2. Inoculate new cultures from scraping of frozen stock into a single tube of Revised Barbour-Stoenner-Kelly Medium.
3. Incubate the tube at 32 to 34°C for 2 to 14 days. Do not shake culture during growth.

Note: Subculturing should be minimized to avoid plasmid loss.<sup>1,7</sup>

**Citation:**

Acknowledgment for publications should read "The following reagent was obtained through BEI Resources, NIAID, NIH: *Borrelia burgdorferi*, Signature-Tagged Mutagenesis Library Clone T03TC121 (Gene IR\_BB\_L35-BB\_L36), NR-22408."

**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](http://www.cdc.gov/biosafety/publications/bmb15/index.htm).

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**References:**

1. Norris, S. J., Personal Communication.
2. Lin, T., et al. "Analysis of an Ordered, Comprehensive STM Mutant Library in Infectious *Borrelia burgdorferi*: Insights into the Genes Required for Mouse Infectivity." *PLoS One* 7 (2012): e47532. PubMed: 23133514.
3. Burgdorfer, W., et al. "Lyme Disease – A Tick-Borne Spirochetosis?" *Science* 216 (1982): 1317-1319. PubMed: 7043737.
4. Johnson, R. C., et al. "*Borrelia burgdorferi* sp. nov.: Etiologic Agent of Lyme Disease." *Int. J. Syst. Bacteriol.* 34 (1984): 496-497.
5. Casjens, S., et al. "A Bacterial Genome in Flux: The Twelve Linear and Nine Circular Extrachromosomal DNAs in an Infectious Isolate of the Lyme Disease Spirochete *Borrelia burgdorferi*." *Mol. Microbiol.* 35 (2000): 490-516. PubMed: 10672174.
6. Fraser, C. M., et al. "Genomic Sequence of a Lyme Disease Spirochaete, *Borrelia burgdorferi*." *Nature* 390 (1997): 580-586. PubMed: 9403685.
7. Purser, J. E. and S. J. Norris. "Correlation Between Plasmid Content and Infectivity in *Borrelia burgdorferi*." *Proc. Natl. Acad. Sci. USA* 97 (2000): 13865-13870. PubMed: 11106398.
8. Kawabata, H., S. J. Norris and H. Watanabe. "BBE02 Disruption Mutants of *Borrelia burgdorferi* B31 Have a Highly Transformable, Infectious Phenotype." *Infect. Immun.* 72 (2004): 7147-7154. PubMed: 15557639.
9. Botkin, D. J., et al. "Identification of Potential Virulence Determinants by *Himar1* Transposition of Infectious *Borrelia burgdorferi* B31." *Infect. Immun.* 74 (2006): 6690-6699. PubMed: 17015459.

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**APPENDIX I: REVISED BSK MEDIUM (ATCC® MEDIUM: 1914)**

HEPES	5.64 g
Neopeptone	4.7 g
Sodium citrate	0.7 g
Glucose	5.64 g
NaHCO <sub>3</sub>	2.0 g
TC-Yeastolate	2.0 g
Sodium pyruvate	0.75 g
N-acetylglucosamine	0.37 g
Bovine serum albumin, fraction V	47.0 g
CMRL 1066, 10X (w/o Glutamine or NaHCO <sub>3</sub> )	100.0 mL
Rabbit serum (heat inactivated)	60.0 mL
Distilled water	840 mL

For agar, add 0.8% agarose.

Dissolve ingredients up to and including bovine serum albumin one at a time in distilled water. Adjust to pH 7.5 with NaOH and filter-sterilize. Aseptically add CMRL 1066 and rabbit serum. Mix well and aseptically dispense into appropriate vessel. Final pH of complete medium should be 7.5 to 7.6.