

# **Certificate of Analysis for NR-51624**

### Genomic DNA from Trypanosoma brucei subsp. gambiense, Strain STIB 386 (in vitro)

#### Catalog No. NR-51624

#### **Product Description:**

Genomic DNA was extracted from *Trypanosoma brucei* (*T. brucei*) subsp. *gambiense*, strain STIB 386 (*in vitro*) was harvested from the blood of infected BALB/c mice and adapted to cell culture by BEI Resources. The parent strain STIB 386 (BEI Resources NR-36198) was derived from strain TH 114/78E (020), which was isolated in 1978 from a male patient in Koudougou, Ivory Coast, West Africa. NR-51624 was extracted from BEI Resources NR-44389 lot 70022602 using proprietary technology. NR-51624 lot 70023641 is provided in 10 mM Tris-HCI, 1 mM EDTA, pH 7.5.

Lot: 70023641 Manufacturing Date: 05FEB2019

TEST	SPECIFICATIONS	RESULTS
Agarose Gel Electrophoresis	High molecular weight chromosomal DNA	High molecular weight chromosomal DNA (Figure 1)
Concentration by PicoGreen® Measurement	0.2 to 3.5 μg in 20 to 200 μL per vial	3 μg in 114 μL per vial (26 μg/mL)
Amount per Vial	0.2 to 3.5 μg	3 µg
Genotypic Analysis Sequencing of internal transcribed spacer (ITS) 1 (1270 base pairs)	≥ 98% sequence identity to <i>T. brucei</i> subsp. <i>gambiense</i> , strain DAL1972 (GenBank: AF306774.1)	98% sequence identity to <i>T. brucei</i> subsp. <i>gambiense</i> , strain DAL1972 (GenBank: AF306774.1) <sup>1</sup>
Functional Activity by PCR Amplification ITS 1, 5.8S ribosomal RNA gene, ITS 2 <sup>2</sup> Serum resistance-associated (SRA) gene <sup>3</sup>	~ 1300 base pair amplicon No amplicon	~ 1300 base pair amplicon No amplicon
OD <sub>260</sub> /OD <sub>280</sub> Ratio	1.7 to 2.1	2
Protozoan Inactivation 10% of total yield inoculated in SDM-79 medium (Life Technologies, custom order part number ME090164 P1) supplemented with 10% heat- inactivated fetal bovine serum and incubated for 14 days at 27°C in an aerobic atmosphere	No viable organisms detected	No viable organisms detected

<sup>&</sup>lt;sup>1</sup>Also consistent with *T. evansi* and/or *T. equiperdum* which are putative subspecies of *T. brucei* (Lun, Z. R., et al. "*Trypanosoma brucei*: Two Steps to Spread Out from Africa." <u>Trends Parasitol</u>. 26 (2010): 424-427. PubMed: 20561822.)

BEI Resources www.beiresources.org E-mail: contact@beiresources.org Tel: 800-359-7370

Fax: 703-365-2898

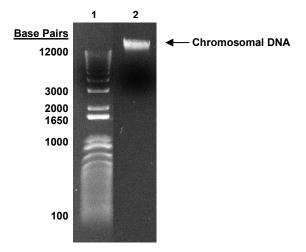
<sup>&</sup>lt;sup>2</sup>PCR was performed as described in Agbo, E. C., et al. "Measure of Molecular Diversity within the *Trypanosoma brucei* Subspecies *Trypanosoma brucei brucei* and *Trypanosoma brucei gambiense* as Revealed by Genotypic Characterization." Exp Parasitol. 99 (2001): 123-131. PubMed: 11846522.

<sup>&</sup>lt;sup>3</sup>T. brucei subsp. gambiense is differentiated from T. brucei subsp. rhodesiense by lack of the SRA gene (Radwanska, M., et al. "The Serum Resistance-Associated Gene as a Diagnostic Tool for the Detection of Trypanosoma brucei rhodesiense." Am. J. Trop. Med. Hyg. 67 (2002): 684-690. PubMed: 12518862.)



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Figure 1: Agarose Gel Electrophoresis



Lane 1: Invitrogen™ TrackIt™ 1 Kb Plus DNA Ladder

Lane 2: ~ 200 ng of NR-51624

/Heather Couch/ Heather Couch

31 MAY 2022

Program Manager or designee, ATCC Federal Solutions

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BEI Resources www.beiresources.org E-mail: contact@beiresources.org
Tel: 800-359-7370

Fax: 703-365-2898