SUPPORTING INFECTIOUS DISEASE RESEARCH

# Toxoplasma gondii, Strain Pru A7 ∆hxgprt::gra2-GFP::tub1-FLUC

## Catalog No. NR-49335

### **Product Description:**

*Toxoplasma gondii (T. gondii)*, strain Pru A7  $\Delta hxgprt::gra2-GFP::tub1-FLUC$  was deposited to BEI Resources as a transgenic strain that expresses green fluorescent protein (GFP) and firefly luciferase (FLUC). Strain Pru A7  $\Delta hxgprt::gra2-GFP::tub1-FLUC$  is derived from strain Prugniaud (Pru)  $\Delta hxgprt$  (also referred to as Pru  $\Delta hpt$ ), which lacks the *hxgprt* (hypoxanthine-xanthine-guanine-phosphoribosyltransferase) gene. The parent strain Prugniaud (Pru) is a Type II strain originally isolated in 1964 from a human with lethal congenital toxoplasmosis in Limoges, France. NR-49335 was produced by cultivation of BEI Resources seed lot 64253082 in human foreskin fibroblast cells (ATCC<sup>®</sup> CRL-1634<sup>TM</sup>) with Dulbecco's Minimal Essential Medium (DMEM) supplemented with 10% (v/v) heat-inactivated fetal bovine serum (HIFBS). The culture was propagated for 3 days at 37°C in an aerobic atmosphere with 5% CO<sub>2</sub> until lysis of the host cell monolayer was reached.

### Lot: 70050592

### Manufacturing Date: 28MAR2022

TEST	SPECIFICATIONS	RESULTS
<b>Cell Morphology<sup>1</sup></b> 7 days at 37°C in an aerobic atmosphere with 5% CO <sub>2</sub> in DMEM supplemented with 10% HIFBS in human	Report result	Refractive; crescent-shaped tachyzoites visible
foreskin fibroblast cells (ATCC <sup>®</sup> CRL-1634™)		
Genotypic Analysis <sup>2</sup> Sequencing of 850 locus (~ 720 base pairs)	≥ 99% sequence identity to <i>T. gondii</i> , strain Prugniaud (GenBank: GU249506.1)	100% sequence identity to <i>T. gondii</i> , strain Prugniaud (GenBank: GU249506.1)
Confirmation of Genes by PCR Amplification <sup>2</sup>		
850 locus <sup>3</sup>	~ 767 base pair amplicon	~ 767 base pair amplicon
850 locus (SfaNI digestion) <sup>3</sup>	Consistent with <i>T. gondii</i> Type II	Consistent with T. gondii Type II
hxgprt locus <sup>4</sup>	No amplicon	No amplicon
Phenotypic Analysis <sup>2</sup>		
Green fluorescent protein (GFP) expression <sup>5</sup>	Positive	Positive
Luciferase activity <sup>6</sup>	Positive	Positive
Viable Cell Count by Hemacytometry <sup>2</sup>	> 10 <sup>6</sup> cells per mL	$6.3 \times 10^7$ cells per mL
Viability <sup>1</sup> 7 days at 37°C in an aerobic atmosphere with 5% CO <sub>2</sub> in DMEM supplemented with 10% HIFBS in human foreskin fibroblast cells (ATCC <sup>®</sup> CRL-1634 <sup>™</sup> )	Growth	Growth
Sterility (21-day incubation) <sup>1</sup>		
Harpo's HTYE broth, 37°C and 26°C, aerobic <sup>7</sup>	No growth	No growth
Trypticase soy broth, 37°C and 26°C, aerobic	No growth	No growth
Sabouraud broth, 37°C and 26°C, aerobic	No growth	No growth
DMEM with 10% FBS, 37°C, aerobic	No growth	No growth
Sheep blood agar, 37°C, aerobic	No growth	No growth
Sheep blood agar, 37°C, anaerobic	No growth	No growth
Thioglycollate broth, 37°C, anaerobic	No growth	No growth
Mycoplasma Contamination DNA Detection by PCR	None detected	None detected

<sup>1</sup>Testing completed on vialed, post-freeze material.

<sup>2</sup>Testing completed on bulk material prior to vialing and freezing.

<sup>3</sup>Primer sequences, annealing temperatures and conditions for restriction enzyme digestion are available at the Toxoplasma Genome Map website (<u>http://toxomap.wustl.edu/Toxo\_Genetic\_Map\_Table.html</u>).

<sup>4</sup>Primer sequences and conditions for PCR are available upon request.

<sup>5</sup>GFP expression was examined by fluorescence microscopy.

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<sup>6</sup>Luciferase activity was determined using the Luciferase Assay System (Promega E1500). Parasites were lysed and incubated with luciferase assay reagent. Light was measured using a spectrophotometer with a wavelength of ~ 560 nm.

<sup>7</sup>Atlas, Ronald M. <u>Handbook of Microbiological Media</u>. 3rd ed. Ed. Lawrence C. Parks. Boca Raton: CRC Press, 2004, p. 798.

#### Figure 1: T. gondii, Strain Pru A7 ∆hxgprt::gra2-GFP::tub1-FLUC – 850 Locus Sequence

CCAGCCGATG CTGCATGGCT GCCACCCCTT CCTCGTAGCC CCCCTGTCGG TGAGGCAACT GGTCCCCGTG GGGTCTTTGA AAGGCTCAAC CGGGTACCCG GGCTACGCGA AGGCGACCC CTTCCACGAG AAGGCCCTC GACCTTATCG ACGCCCGTTC GCCCTGGCGG CGGCTTGCCA TCGCTTCTGG ATGTCGGCGC TGCTTTCCCT GGAGGCACCC CTGTTTGTG GGAGGAAGCA GTAGTGGTAC TAATGGGTGC CTGTGCCTGC CTCCCTCCTT GCTCCGGTGA GCTGCTGCCC CCCACAGGTC CCTCTTGCTG CATGCCTCGT GCAGGAGCGC TTGGCGTCGG TGAGTCACCA TCTGACGGTG AAACTGAGCT GTCGCCCAAG CCGCTGCTGC CTGACGACGA AGAAGGCCCA CCTGAGTGGA TTATGACGAC GTCACCCTCT GGCCCGCTTT CAGAGCCCAC GAAAAATGAA GCAAGACGCG GGCCTTGCA TGGGGACGGT GATGGCGGCG AACGTTTTCC GGGAACGTG GTTCCGATGT CTCTTGTCGG GGATGTTTCC TTTGGAAGCG AGCCCTTTGC GCCGCAGCAC GGACTTTGTG TGTCTGCTGG TACAAGGACA GTCTCTGAAG GACTCCCCGT TGCAGGTGTG GAATGTAAAG GCCCCTCTAC GGGTTTCCC CCAGATGCTG GAGAAGGTGG TGATCTGGAG GCCGAAACAT

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