

Plasmodium falciparum, Strain NF54::DiCre

Catalog No. MRA-1314

Product Description:

Plasmodium falciparum (*P. falciparum*), strain NF54::DiCre is a derivative of the NF54 strain with the rapamycin-induced dimerized Cre recombinase (DiCre) system integrated into the *pf547* locus via CRISPR/Cas9. MRA-1314 was produced by cultivation of the deposited material in fresh human erythrocytes suspended in RPMI 1640 medium, adjusted to contain 10% (v/v) heat-inactivated human serum (pooled Type A+), 25 mM HEPES, 2 mM L-glutamine, 4 g/L D-glucose, 0.005 µg/mL hypoxanthine and 2.5 µg/mL gentamicin. The culture was incubated at 37°C in sealed flasks outgassed with blood-gas atmosphere (90% N₂, 5% CO₂, 5% O₂) and monitored for parasitemia every 1 to 4 days for 17 days. Every 1 to 4 days, uninfected, leukocyte filtered, Type O erythrocytes in complete culture medium were added dropwise to the culture as needed and monitored for hematocrit.

Lot: 70028492

Manufacturing Date: 27SEP2019

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TEST	SPECIFICATIONS	RESULTS
Identification by Giemsa Stain Microscopy¹	Blood-stage parasites present	Blood-stage parasites present
Antimalarial Susceptibility Profile (<i>in vitro</i>)¹ Half-maximal Inhibitory Concentration (IC ₅₀) by SYBR green I [®] drug sensitivity assay ² Chloroquine Artemisinin Quinine Cycloguanil Pyrimethamine Sulfadoxine	Report results Report results Report results Report results Report results Report results	10.5 ± 0.5 nM 10.5 ± 0.2 nM 51.2 ± 2.4 nM 15.4 ± 1.1 nM 55.5 ± 3.8 nM 410900 ± 28407 nM
Genotypic Analysis³ Sequencing of Merozoite Surface Protein 2 (MSP2) gene (~ 800 base pairs)	≥ 99% sequence identity to <i>P. falciparum</i> , strain NF54::DiCre (GenBank: QFXU01000003.1)	100% sequence identity to <i>P. falciparum</i> , strain NF54::DiCre (GenBank: QFXU01000003.1) (Figure 1)
Functional Activity by PCR Amplification³ MSP2 PCR amplicon analysis	~ 600-900 base pair amplicon	~ 900 base pair amplicon
Level of Parasitemia by Giemsa Stain Microscopy Pre-freeze (17 days post-infection) ³ Ring-stage parasitemia Total parasitemia Post-freeze (3 days post-infection) ¹ Ring-stage parasitemia Total parasitemia	Report results ≥ 2% Report results ≥ 1%	2.54% 4.76% 0.55% 1.84%
Viability (post-freeze; 3 days post-infection)¹	Growth in infected red blood cells	Growth in infected red blood cells
Sterility (21-day incubation)¹ Harpo's HTYE broth, 37°C and 26°C, aerobic ⁴ Trypticase soy broth, 37°C and 26°C, aerobic Sabouraud broth, 37°C and 26°C, aerobic DMEM with 10% FBS, 37°C, aerobic Sheep blood agar, 37°C, aerobic Sheep blood agar, 37°C, anaerobic Thioglycollate broth, 37°C, anaerobic	No growth No growth No growth No growth No growth No growth No growth	No growth No growth No growth No growth No growth No growth No growth

TEST	SPECIFICATIONS	RESULTS
Mycoplasma Contamination¹ DNA detection by PCR	None detected	None detected

¹Testing completed on vial, post-freeze material.

²A SYBR Green I[®] anti-malarial drug sensitivity assay in 96-well plates was used to determine IC₅₀ values of an active (> 70% ring stage) parasite culture in the presence of each antimalarial drug [Hartwig, C. L., et al. "XI: I. SYBR Green I[®]-Based Parasite Growth Inhibition Assay for Measurement of Antimalarial Drug Susceptibility in *Plasmodium falciparum*." In *Methods in Malaria Research Sixth Edition*. (2013) Moll, K., et al. (Ed.), EVIMalaR, pp. 122-129. Available at: <https://www.beiresources.org/Publications/MethodsInMalariaResearch.aspx>.]

³Testing completed on bulk material prior to vialing and freezing.

⁴Atlas, Ronald M. *Handbook of Microbiological Media*. 3rd ed. Ed. Lawrence C. Parks. Boca Raton: CRC Press, 2004, p. 798.

Figure 1: MRA-1314 MSP2 Sequence

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AATTAAAACA TTGTCTATTA TAAATTTCTT TATTTTTGTT ACCTTTAATA TTAAAAATGA AAGTAAATAT AGCAACACAT
TCATAAACAA TGCTTATAAT ATGAGTATAA GGAGAAGTAT GGCAGAAAGT AAGCCTTCTA CTGGTGCTGG TGGTAGTGCT
GGTGGTAGTG CTGGTGGTAG TGCTGGTGGT AGTGCTGGTG GTAGTGCTGG TGGTAGTGCT GGTTCTGGTG ATGGTAATGG
TGCAGATGCT GAGGGAAGTT CAAGTACTCC CGCTACTACC ACAACTACCA AAACCTACCAC AACTACCACA ACTACTAATG
ATGCAGAAGC ATCTACCAGT ACCTCTTCAG AAAATCCAAA TCATAAAAAAT GCCGAAACAA ATCCAAAAGG TAAAGGAGAA
GTTCAAGAAC CAAATCAAGC AAATAAAGAA ACTCAAAATA ACTCAAATGT TCAACAAGAC TCTCAAATA AATCAAATGT
TCCACCCACT CAAGATGCAG AACTAAAAG TCCTACTGCA CAACCTGAAC AAGCTGAAAA TTCTGCTCCA ACAGCCGAAC
AAACTGAATC CCCCGAATTA CAATCTGCAC CAGAGAATAA AGGTACAGGA CAACATGGAC ATATGCATGG TTCTAGAAAT
AATCATCCAC AAAATACTTC TGATAGTCAA AAAGAATGTA CCGATGGTAA CAAAGAAAAC TGTGGAGCAG CAACATCCCT
CTTAAATAAC TCTAGTAATA TTGCTTCAAT AAATAAATTT GTTGTTTTAA TTTCAGCAAC ACTTGTTTTA TCTTTTGCCA
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