

***Plasmodium falciparum*, Strain CamWT\_C580Y**

**Catalog No. MRA-1251**

**Product Description:** *Plasmodium falciparum* (*P. falciparum*), strain CamWT\_C580Y is a K13-propeller mutant of the CamWT strain (BEI Resources MRA-1250), featuring a single nucleotide substitution leading to a C580Y amino acid change. *P. falciparum*, strain CamWT\_C580Y was deposited as more resistant to artemisinin than the parent strain, with a ring-stage survival assay (RSA<sub>0-3h</sub>) value of 8.9% when exposed to dihydroartemisinin.

**Lot<sup>1</sup>: 70018495**

**Manufacturing Date: 27SEP2018**

TEST	SPECIFICATIONS	RESULTS
<b>Identification by Giemsa Stain Microscopy<sup>2,3</sup></b>	Blood-stage parasites present	Blood-stage parasites present
<b>Antimalarial Susceptibility Profile (<i>in vitro</i>)<sup>2</sup></b> Half-maximal Inhibitory Concentration (IC <sub>50</sub> ) by SYBR green I <sup>®</sup> drug sensitivity assay <sup>4</sup> Chloroquine Artemisinin Quinine Cycloguanil Pyrimethamine Sulfadoxine Ring-stage Survival Assay (RSA <sub>0-3h</sub> ) <sup>5</sup> Dihydroartemisinin (DHA) <sup>6</sup>	Report results Report results Report results Report results Report results Report results Report results	45.8 ± 6.3 nM 12.2 ± 0.8 nM 68.7 ± 6.3 nM 519 ± 23.9 nM 19740 ± 2278 nM 409500 ± 37770 nM 9.44%
<b>Genotypic Analysis<sup>2</sup></b> Sequencing of Merozoite Surface Protein 2 (MSP2) gene (~ 750 base pairs) Sequencing of kelch protein (K13-propeller) gene <sup>7</sup> (~ 1990 base pairs)	Consistent with <i>P. falciparum</i> Contains K13 C580Y mutation	Consistent with <i>P. falciparum</i> (Figure 1) Contains K13 C580Y mutation (Figure 2)
<b>Functional Activity by PCR Amplification<sup>2</sup></b> MSP2 PCR amplicon analysis <sup>8</sup>	~ 600 to 900 base pair amplicon	~ 900 base pair amplicon
<b>Level of Parasitemia</b> Pre-freeze <sup>9,10</sup> Ring-stage parasitemia Total parasitemia Post-freeze <sup>2,11</sup> Ring-stage parasitemia Total parasitemia	Report results ≥ 2%  Report results ≥ 1%	3.58% 5.07%  1.26% 3.78%
<b>Viability<sup>2,12</sup></b>	Growth in infected red blood cells	Growth in infected red blood cells
<b>Sterility (21-day incubation)<sup>2</sup></b> Harpo's HTYE broth <sup>13</sup> , 37°C and 26°C, aerobic Tryptic Soy broth, 37°C and 26°C, aerobic Sabouraud Dextrose 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
<b>Mycoplasma Contamination<sup>2</sup></b> DNA Detection by PCR	None detected	None detected

<sup>1</sup>MRA-1251 was produced by cultivation of BEI Resources MR-MRA-1251 lot 62368019 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% N<sub>2</sub>, 5% CO<sub>2</sub>, 5% O<sub>2</sub>) and monitored for parasitemia daily for 22 days. Every 1 to 3 days, uninfected, leukocyte filtered, Type O erythrocytes in complete culture medium were added dropwise to the culture as needed and monitored for hematocrit.

<sup>2</sup>Testing completed on vial post-freeze material

<sup>3</sup>Blood-stage malaria parasites (rings, trophozoites, schizonts +/- gametocytes) were examined by microscopic Giemsa-stained blood smears of an *in vitro* human blood culture over 5 days.

<sup>4</sup>A SYBR Green I<sup>®</sup> anti-malarial drug sensitivity assay in 96-well plates was used to determine IC<sub>50</sub> 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<sup>®</sup>-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>].

<sup>5</sup>A detailed RSA<sub>0-3h</sub> protocol is available on the Worldwide Antimalarial Resistance Network's website at <http://www.wwarn.org/tools-resources/procedures/ring-stage-survival-assays-rsa-evaluate-vitro-and-ex-vivo-susceptibility>.

<sup>6</sup>*P. falciparum*, strain CamWT\_C580Y was reported with a DHA RSA<sub>0-3h</sub> value of 8.9% [Straimer, J., et al. "Drug Resistance. K13-Propeller Mutations Confer Artemisinin Resistance in *Plasmodium falciparum* Clinical Isolates." *Science* 347 (2015): 428-431. PubMed: 25502314.].

<sup>7</sup>K13-propeller mutation C580Y confers artemisinin resistance *in vitro*; for additional information, please refer to Straimer, J., et al. "Drug Resistance. K13-Propeller Mutations Confer Artemisinin Resistance in *Plasmodium falciparum* Clinical Isolates." *Science* 347 (2015): 428-431. PubMed: 25502314.

<sup>8</sup>Primer sequences and conditions for PCR are available upon request.

<sup>9</sup>Testing completed on bulk material prior to vialing and freezing

<sup>10</sup>Parasitemia was determined after 22 days post infection by microscopic counts of Giemsa-stained blood smears.

<sup>11</sup>Parasitemia was determined after 5 days post infection by microscopic counts of Giemsa-stained blood smears.

<sup>12</sup>Viability was confirmed by examination of infected erythrocytes for parasitemia at 5 days post infection.

<sup>13</sup>Atlas, Ronald M. *Handbook of Microbiological Media*. 3rd ed. Ed. Lawrence C. Parks. Boca Raton: CRC Press, 2004, p. 798.

**Figure 1: MRA-1251 MSP2 Sequence**

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AAACATTGTC TATTATAAAT TTCTTTATTT TTGTTACCTT TAATATTAAA AATGAAAGTA AATATAGCAA CACATTCATA
AACAAATGCTT ATAATATGAG TATAAGGAGA AGTATGGCAA ATGAAGGTTT TAATACTAAT AGGGTAGGTG CAAATGCTCC
AAAAGCTGAT ACTATTGCTA GTGGAAGTCA AAGTAGTACA AATAGTGCAA GTACTAGTAC TACTAATAAT GGAGAATCAC
AACTACTAC TCCTACCGCT GCTGATACCC CTACTGCTAC AAAAAGTAAT TCACCTTCAC CACCCATCAC TACTACAGAA
AGTAATTCAC CTTACACCAC CATCACTACT ACAGAAAGTA ATTCACCTTC ACCACCCATC ACTACTACAG AAAGTTCAAG
TTCTGGCAAT GCACCAAATA AAACAGACGG TAAAGGAGAA GAGAGTGAAA AACAAAATGA ATTAAATGAA TCAACTGAAG
AAGGACCCAA AGCTCCACAA GAACCTCAAA CGGCAGAAAA TGAAAATCCT GCTGCACCAG AGAATAAAGG TACAGGACAA
CATGGACATA TGCATGGTTC TAGAAATAAT CATCCACAAA ATACTTCTGA TAGTCAAAAA GAATGTACCG ATGGTAACAA
AGAAAATGT GGAGCAGCAA CATCCCTCTT AAATAACTCT AGTAATATTG CTTCAATAAA TAAATTTGTT GTTTTAAATTT
CAGCAACACT TGTTTTATCT TTTGCCATA
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**Figure 2: MRA-1251 K13 Sequence**

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GGGAACATAA AAAGATGGGC CAAGCTGCCA TTCATTTGTA TCTGGTGAAA AGAAATGACA TGAATTTAGA ACTTCGCCAT
TTTCTCCTCC TGTAATTATA TAAGAATCTG ACAATGTGGC AGCTCCAAAA TTCATTTTTT TCTCTGGTAC ACCATTTAGA
AATTGCCATC TTTTATTAAA TGTTTGATAT TGTTCAACGG AATCTAATAT GTTATGTTCA TTATCAATAC CTTCAACAAC
ATATATTTGA TTAAGGTAAT TAAAAGCTGC TCCTGAACCT CTAGCTTCTA ATAAGGCATA TGGAAATGTT TCCCATTTAT
TCATTTTTTC TTCATATACT TCAATAGAAT TTAATCTCTC ACCATTAGTT CCACCAATGA CATAAATTTT ATTATCAAAA
GCAACATACA TAGCTGATGA TCTAGGGGTA TTCAAAGGTG CCACCTCTAC CCATGCTTTC ATACGATGAT CATATGCTTC
TACATTCGGT ATAATAGAAG AGCCATCATA TCCCCAATA CAATAAATTC TACCATTTGA CGTAACACCA CAATTATTTT
TTCTAGGTAT ATTTAAATTA CTTGATACAT ACCAGACGTC TCTTAATCTA TCATACACCT CAGTTTCAAA TAAAGCCTTA
TAATCATAGT TATTACCACC AAAAACGTAT AAGAAATTAT TCAATACAGC ACTTCCAAAA TAAGCTTTTT TGGTAGACAT
AGGTGTACAC ATACGCCAGC ATTGTTGACT AATATCTAAT AATTCCATCG AATTTAAATA TTCTACACCA TCAAATCCAC
CTATACAAAA TACTAATGGG AATGGTAAAA ATTTAATACC ATAAAATCT GCTTCTTTCA ACAAGGCTTC ACTTTCCTT
AAATCTTTTG GTATGGGTAT AGTTAACGGA TTTCTTAAGA AGTTAAGTAT AATTCTAAAT AACTCACTAT CCCTATCTAA
GAATATTCTT CCTTGTTTAT CTCTGGTTAC ATGATGTCTT CCACTTAATA ATTTCTCTAT AAATGAATCT TTTTGTTGTG
TTAAGGTATG TCTAGATGTT TCAAAAATAG CTCCACCAAC ATTAATATCA ATCATAGTTT CAGTAGCAAT ATTTGCATCA
ACAATTTTTT TTTTATCATT CTCAAAATTT TTATATTCTT GATAATATTT TTCTTTTTCT AAATATAATA CTAATTTAAT
TTTATCGATT TCTTGTAATA ATCTTAATCT TTCTTCATCA AATCGTTTCC TATGTTCTTC TTTTCTTTT TTTATTTGTT
TATAACCATT AGATATATCA ATATCTAATT TCTTTCTTTC ATCATGTAAT TTCTGTTCTT CAATATTTTT ACGGTTTTCT
AATTCTTTGT ACAATCGTAC TCTTTCCATT TCTAGTTCTT TCTTATCTTT AAATAATTTA TCTTTTTCTC GAATAAAATT
CATTTGTGTC TTTTTTAACC AATTAATAAA TGTAATTCTT AAATCACCTA CCATATTTTC AAAATCACTA GCATCACTTA
ATTCCGTTTC AATAATTTTC TTTTCATAAG TATCATTTAC ATTATTTACA GTTGAAAATT TTTTAAATTT GTTTAATCCA
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GAATCATCAT TTATAAGATT TGCTGCATCT AAAAAATTCT CTTTTTTGTT GGTATTCATA ATTGATGGAG AATTCATATT  
 ATTATTAATA AGATTATTAG TTATATTATT TGCTGTCAGA TTATTATTAT TATTATTATT ATTATGATTA ATATTATTAT  
 TTTTCATTTGT AATAGTATCT TTTTTTTTAT TTAACAATTT ATTTAAATAT TTATTTCCCTA TATTATCTTT AGACATATTA  
 TTAAATGTTT TTAGATAAATT ACTTGGTAAA AAATCTTTTT TACTATCAAA GTTCGAATCT AATACACTCA TATCAATGGA  
 TTCTAATAGG CTATCTTTAA CATTTCATA ACTACTATTA TTTAAAAGGA AACTATTATT TTCCGTTTTT TC

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