

# Copy number estimation of *P. falciparum* pfmdr1 v1.1

## Procedure



**Molecular Module**

**WorldWide Antimalarial Resistance Network (WWARN)**



**Suggested citation:** Molecular Module. 2011. Copy number estimation of *P. falciparum* pfmdr1 v1.1. WWARN Procedure

**Procedure ID:** [Insert unique ID– to be generated from a master list]

**This procedure was developed by:**

Ric Price, Menzies School of Health Research, Australia; WWARN

Molecular Module, WWARN

#### Version History

Version number	Revision(s) & reason for amendment	Release date
1.1	Change from 50 ng/uL template DNA to 50 pg/uL	17/03/2011
1.0	Creation of procedure	18/11/2010

**For more information, contact:**

molecular@wwarn.org

WorldWide Antimalarial Resistance Network (WWARN)

[www.wwarn.org](http://www.wwarn.org)

## Contents

1. Purpose .....	4
2. Scope .....	4
3. Abbreviations .....	4
4. Duties and Responsibilities .....	4
5. Materials and Equipment .....	4
5.1 Materials .....	4
Table 1. Commonly used pfmdr1 copy number control strains .....	5
5.2 Equipment.....	5
6. Procedure.....	5
6.1 Real-time multiplex PCR .....	5
Table 2. Real-time multiplex PCR master mix.....	6
Table 3. Real-time PCR thermal cycling conditions .....	7
6.2 Quality check and estimation of copy number .....	7
7. References .....	7

## 1. Purpose

This procedure is designed to estimate the copy number of the *Plasmodium falciparum* multi-drug resistance 1 locus (pfmdr1) in DNA extracted from samples of infected blood.

## 2. Scope

This procedure is intended for use in clinical trials and molecular studies where whole blood or dried blood spot samples are collected for genotyping of *P. falciparum* infections. It describes the relative quantification of the pfmdr1 gene associated with resistance to multiple antimalarials. This procedure is applicable for well-equipped laboratories familiar with real-time PCR.

## 3. Abbreviations

DNA	Deoxyribonucleic Acid
pfmdr1	<i>Plasodium falciparum</i> multi-drug resistance 1 locus
PCR	Polymerase Chain Reaction
µL	microlitre
mL	millilitre
pg	picogram
nM	nanomolar
ΔΔ C <sub>T</sub>	Delta delta Cycle threshold
C <sub>T</sub>	Cycle threshold

## 4. Duties and Responsibilities

N/A

## 5. Materials and Equipment

### 5.1 Materials

- 2X TaqMan Master Mix (ABI, part # 4304437) or comparable reagent
- MicroAMP Optical 96-well plates (ABI, part # N801-0560) or comparable optical PCR tubes/plates

- MicroAMP Optical Adhesive Film Covers (ABI, part # 4311971) or comparable optical PCR film/caps
- Micropipets and tips (10  $\mu$ L, 200  $\mu$ L and 1000  $\mu$ L)
- 1.5 mL microcentrifuge tubes or 15 mL conical tubes for master mix (depending on number of samples)
- pfmdr1 primers and probe (target locus):
  - F: 5'-TGC ATC TAT AAA ACG ATC AGA CAA A-'3
  - R: 5'-TCG TGT GTT CCA TGT GAC TGT-'3
  - Probe: 5'-6FAM-TTT AAT AAC CCT GAT CGA AAT GGA ACC TTT G-TAMRA-'3
- $\beta$ -tubulin primers and probe (endogenous locus):
  - F: 5'-AAA AAT ATG ATG TGC GCA AGT GA-'3
  - R: 5'-AAC TTC CTT TGT GGA CAT TCT TCC T-'3
  - Probe: 5'-VIC-TAG CAC ATG CCG TTA AAT ATC TTC CAT GTC T-TAMRA-'3
- DNA controls: at least one single-copy template and one multi-copy template (see Table 1) at 50 pg/ $\mu$ L.

Table 1. Commonly used pfmdr1 copy number control strains

<u>Strain</u>	<u>Approx. number of copies</u>
3D7	1
K1	1
D10	1
Fac8	2-3
Dd2	3-4

## 5.2 Equipment

- Real-Time PCR machine

## 6. Procedure

### 6.1 Real-time multiplex PCR

Real-time PCR can be performed using a number of real-time PCR machines and reagents. The procedure given here uses the ABI system and probes with 5'-6FAM or VIC fluorescent labels with a 3'-TAMRA quencher for detection. Other machines and reagents, such as Minor Groove Binding (MGB)-labeled primers<sup>1</sup>, require different reaction mixes, sample volumes and/or cycling conditions.

Success of this assay may vary based on parasitaemia and quality of the sample, particularly when using dried blood spots. Greater success is generally achieved using DNA extracted from samples with > 10,000 parasites/ $\mu$ L<sup>2</sup>.

- I. Calculate the number of samples to be tested. Each control and sample is quantified in triplicate multiplex reactions for *pfmdr1* and  $\beta$ -tubulin. A representative 96-well plate is given in Figure 1.

	1	2	3	4	5	6	7	8	9	10	11	12
A	3D7	3D7	3D7	5	5	5	13	13	13	21	21	21
B	K1	K1	K1	6	6	6	14	14	14	22	22	22
C	Dd2	Dd2	Dd2	7	7	7	15	15	15	23	23	23
D	water	water	water	8	8	8	16	16	16	24	24	24
E	1	1	1	9	9	9	17	17	17	25	25	25
F	2	2	2	10	10	10	18	18	18	26	26	26
G	3	3	3	11	11	11	19	19	19	27	27	27
H	4	4	4	12	12	12	20	20	20	28	28	28

*pfmdr1* +  $\beta$ -tubulin

Figure 1. 96-well plate including template controls, a no-template control and 28 samples in triplicate multiplexed reactions. Image source: Meera Venkatesan, University of Maryland Baltimore.

- II. Place a 1.5 mL or 15 mL tube on ice. Prepare PCR master mix by adding 2x TaqMan Master Mix reagent, forward and reverse primers and probes in the amounts given in Table 2. It is best to prepare the master mix for 2–4 extra samples so that it does not run out.

Table 2. Real-time multiplex PCR master mix

<u>Reagent</u>	<u>Amount per reaction</u>
2x TaqMan Master Mix reagent	12.5 $\mu$ L
<i>Pfmdr1</i> Forward primer	300 nM
<i>Pfmdr1</i> Reverse primer	300 nM
<i>Pfmdr1</i> probe	100 nM
$\beta$ -tubulin Forward primer	300 nM
$\beta$ -tubulin Reverse primer	300 nM
$\beta$ -tubulin probe	100 nM
Water	to reach total of 20 $\mu$ L

**NOTE:** Do not vortex the mix. Mix by hand-shaking and light centrifugation. Avoid prolonged exposure of mix to light.

- III. Add 20  $\mu$ L of master mix to each well. Add 5  $\mu$ L of control DNA, water or sample DNA to each well as shown in Figure 1.
- IV. Seal plate thoroughly and cycle under the PCR conditions listed in Table 3. Label the single-copy control (usually 3D7) as the calibrator.

Table 3. Real-time PCR thermal cycling conditions

Step	Temperature (°C)	Time
1	95	5:00
2	95	0:15
3*	58	0:60
4	Repeat steps 2–3 40 times	

\*Data collection occurs in step 3 shown in **green**

## 6.2 Quality check and estimation of copy number

- I. Check the quality of your reaction. Repeat sample or control if<sup>2</sup>:
  - a. The spread of  $\Delta\Delta C_T$  values among replicates exceeds 1.5.
  - b.  $C_T$  estimate > 35.
  - c. Copy number estimate ranges between 1.3–1.6.

**NOTE:** Other quality cutoff rules are also commonly used, including repetition or exclusion of samples with a  $C_T$  estimate > 32, with an upper limit of the 95% confidence interval of the copy number estimate > 0.4 or a copy number estimate of < 0.5<sup>3,4</sup>.

## 7. References

1. Nair S, Nash D, Sudimack D, Jaidee A, Barends M, Uhlemann AC, Krishna S, Nosten F, Anderson TJ. 2007. **Recurrent gene amplification and soft selective sweeps during evolution of multidrug resistance in malaria parasites.** Mol Biol Evol. 24:562-573.
2. Price RN, Uhlemann AC, Brockman A, McGready R, Ashley E, Phaipun L, Patel R, Laing K, Looareesuwan S, White NJ, Nosten F, Krishna S. 2004. **Mefloquine resistance in Plasmodium falciparum and increased pfmdr1 gene copy number.** Lancet. 364:438-447.
3. Nair S, Miller B, Barends M, Jaidee A, Patel J, Mayxay M, Newton P, Nosten F, Ferdig MT, Anderson TJ. 2008. **Adaptive Copy Number Evolution in Malaria Parasites.** PLoS Genet. 4: e1000243. doi:10.1371/journal.pgen.1000243
4. Griffing S, Syphard L, Sridaran S, McCollum AM, Mixson-Hayden T, Vinayak S, Villegas L, Barnwell JW, Escalante AA, Udhayakumar V. 2010. **pfmdr1 amplification and fixation of pfcr1 chloroquine resistance alleles in Plasmodium falciparum in Venezuela.** Antimicrob Agents Chemother. 54:1572-1579.