

Documents

Yusuf, N.M.^a, Azman, A.N.^{a b}, Aziz, A.A.A.^{a c}, Fuad, F.A.A.^b, Nasarudin, R.N.^a, Hisam, S.^a

Evaluation of the binding interactions between Plasmodium falciparum Kelch-13 mutant recombinant proteins with artemisinin

(2024) *PLoS ONE*, 19 (8 August), art. no. e0306975, .

DOI: 10.1371/journal.pone.0306975

^a Parasitology Unit, Infectious Disease Research Centre, Institute for Medical Research, National Institute of Health, Shah Alam, Malaysia

^b Department of Chemical Engineering & Sustainability, Faculty of Engineering, International Islamic University Malaysia, Kuala Lumpur, Malaysia

^c School of Biology, Faculty of Applied Sciences, Universiti Teknologi MARA (UiTM) Cawangan Negeri Sembilan, Kampus Kuala Pilah, Kuala Pilah, Malaysia

Abstract

Malaria, an ancient mosquito-borne illness caused by Plasmodium parasites, is mostly treated with Artemisinin Combination Therapy (ACT). However, Single Nucleotide Polymorphisms (SNPs) mutations in the P. falciparum Kelch 13 (PfK13) protein have been associated with artemisinin resistance (ART-R). Therefore, this study aims to generate PfK13 recombinant proteins incorporating of two specific SNPs mutations, PfK13-V494I and PfK13-N537I, and subsequently analyze their binding interactions with artemisinin (ART). The recombinant proteins of PfK13 mutations and the Wild Type (WT) variant were expressed utilizing a standard protein expression protocol with modifications and subsequently purified via IMAC and confirmed with SDS-PAGE analysis and Orbitrap tandem mass spectrometry. The binding interactions between PfK13-V494I and PfK13-N537I propeller domain proteins ART were assessed through Isothermal Titration Calorimetry (ITC) and subsequently validated using fluorescence spectrometry. The protein concentrations obtained were 0.3 mg/ml for PfK13-WT, 0.18 mg/ml for PfK13-V494I, and 0.28 mg/ml for PfK13-N537I. Results obtained for binding interaction revealed an increased fluorescence intensity in the mutants PfK13-N537I (83 a.u.) and PfK13-V494I (143 a.u.) compared to PfK13-WT (33 a.u.), indicating increased exposure of surface proteins because of the looser binding between PfK13 protein mutants with ART. This shows that the PfK13 mutations may induce alterations in the binding interaction with ART, potentially leading to reduced effectiveness of ART and ultimately contributing to ART-R. However, this study only elucidated one facet of the contributing factors that could serve as potential indicators for ART-R and further investigation should be pursued in the future to comprehensively explore this complex mechanism of ART-R. © 2024 Md. Yusuf et al. This is an open access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

Index Keywords

artemisinin, Plasmodium falciparum kelch-13, recombinant protein, unclassified drug, antimalarial agent, artemisinin, artemisinin derivative, protein binding, protozoal protein; Article, cloning vector, documentation, evaluation study, gene expression, gene mutation, gene sequence, immobilized metal affinity chromatography, isothermal titration calorimetry, ligand binding, liquid chromatography-mass spectrometry, mass spectrometry, nonhuman, protein expression, protein interaction, protein purification, real time polymerase chain reaction, sequence analysis, signal detection, single nucleotide polymorphism, spectrofluorometry, chemistry, drug effect, drug resistance, genetics, metabolism, mutation, Plasmodium falciparum; Antimalarials, Artemisinins, Drug Resistance, Mutation, Plasmodium falciparum, Polymorphism, Single Nucleotide, Protein Binding, Protozoan Proteins, Recombinant Proteins

Chemicals/CAS

artemisinin, 63968-64-9; Antimalarials; artemisinin; Artemisinins; Protozoan Proteins; Recombinant Proteins

Manufacturers

Agilent; Thermo

References

- (2019) *Report on antimalarial drug efficacy, resistance and response*, 63–1,2,3,4,5,6
- Amambua-Ngwa, A, Button-Simons, KA, Li, X, Kumar, S, Brenneman, KV, Ferrari, M **Chloroquine resistance evolution in Plasmodium falciparum is mediated by the putative amino acid transporter AAT1** (2023) *Nat Microbiol*, 8 (7), pp. 1213-1226. PMID: 37169919
- Sibley, CH, Hyde, JE, Sims, PFG, Plowe, C V, Kublin, JG, Mberu, EK **Pyrimethamine–sulfadoxine resistance in Plasmodium falciparum: what next?** (2001) *Trends Parasitol*, 17 (12), pp. 570-571.

- Sidhu, ABS, Verdier-Pinard, D, Fidock, DA.
Chloroquine Resistance in Plasmodium falciparum Malaria Parasites Conferred by pfcr Mutations
(2002) *NIH Public Access*, 298 (5591), pp. 210-213.
- Ridley, RG.
Malaria: Dissecting chloroquine resistance
(1998) *Current Biology*, 8 (10), pp. 346-349.
PMID: 9601634
- Guémas, E, Coppée, R, Ménard, S, du Manoir, M, Nsango, S, Makaba Mvumbi, D
Evolution and spread of Plasmodium falciparum mutations associated with resistance to sulfadoxine–pyrimethamine in central Africa: a cross-sectional study
(2023) *Lancet Microbe*, 4 (12), pp. e983-e993.
- Van Lenthe, M, Van Der Meulen, R, Lassoovski, M, Ouabo, A, Bakula, E, Badio, C
Markers of sulfadoxine-pyrimethamine resistance in Eastern Democratic Republic of Congo; Implications for malaria chemoprevention
(2019) *Malar J*, 18 (1), pp. 1-9.
[Internet]
- Mbugi, E V., Mutayoba, BM, Malisa, AL, Balthazary, ST, Nyambo, TB
Mshinda. Drug resistance to sulphadoxine-pyrimethamine in Plasmodium falciparum malaria in Mlimba, Tanzania
(2006) *Malar J*, 5, pp. 1-11.
- Lin, JT, Juliano, JJ, Wongsrichanalai, C.
Drug-resistant malaria: The era of ACT
(2010) *Curr Infect Dis Rep*, 12 (3), pp. 165-173.
PMID: 21308525
- (2021) *World malaria report 2021*,
[Internet] [Internet]. World Health Organization. 2013–2015
- Miller, LH, Su, X.
Artemisinin: Discovery from the Chinese herbal garden
(2011) *Cell*, 146 (6), pp. 855-858.
[Internet]. PMID: 21907397
- **Management Guideline for Malaria in Malaysia**
(2014) *Cpg*, 59.
[Internet]. ;. https://www.google.com/url?sa=t&rct=j&q=&esrc=s&source=web&cd=&cad=rja&uact=8&ved=2ahUKEwjatLmJxPP4AhVFAGwGHXL-D4oQFnoECAcQAQ&url=https%3A%2F2Fwww.moh.gov.my%2Findex.php%2Ffile_manager%2Fdl_item%2F554756755a584a69615852686269394859584a70637942515957356b6
- Noedl, H, Se, Y, Schaecher, K, Smith, BL, Socheat, D, Fukuda, MM.
Evidence of Artemisinin-Resistant Malaria in Western Cambodia
(2008) *New England Journal of Medicine*, 359 (24), pp. 2619-2620.
[Internet]. Dec 11 [cited 2019 Apr 18]; PMID: 19064625
- Dondorp, AM, Nosten, F, Yi, P, Das, D, Phyto, AP, Tarning, J
Artemisinin Resistance in
(2009) *Drug Ther (NY)*, 361 (5), pp. 455-467.
[Internet]
- Tun, KM, Imwong, M, Lwin, KM, Win, AA, Hlaing, TM, Hlaing, T
Spread of artemisinin-resistant Plasmodium falciparum in Myanmar: a cross-sectional survey of the K13 molecular marker
(2015) *Lancet Infect Dis*, 15 (4), pp. 415-421.
[Internet]. Apr [cited 2019 Apr 9]
- Bonnington, CA, Phyto, AP, Ashley, EA, Imwong, M, Sriprawat, K, Parker, DM
Plasmodium falciparum Kelch 13 mutations and treatment response in patients in Hpa-Pun

- District, Northern Kayin State, Myanmar**
(2017) *Malar J*, 16 (1), pp. 1-7.
[Internet]
- Phyo, AP, Nkhoma, S, Stepniewska, K, Ashley, EA, Nair, S, McGready, R
Emergence of artemisinin-resistant malaria on the western border of Thailand: a longitudinal study
(2012) *The Lancet*, 379 (9830), pp. 1960-1966.
[Internet]. May 26 [cited 2019 Apr 9]; PMID: 22484134
 - Amaratunga, C, Sreng, S, Suon, S, Phelps, ES, Stepniewska, K, Lim, P
Artemisinin-resistant Plasmodium falciparum in Pursat province, western Cambodia: a parasite clearance rate study
(2012) *Lancet Infect Dis*, 12 (11), pp. 851-858.
[Internet]. Nov 1 [cited 2019 Apr 9]; [https://doi.org/10.1016/S1473-3099\(12\)70181-0](https://doi.org/10.1016/S1473-3099(12)70181-0) PMID: 22940027
 - Miotto, O, Sekihara, M, Tachibana, SI, Yamauchi, M, Pearson, RD, Amato, R
Emergence of artemisinin-resistant Plasmodium falciparum with kelch13 C580Y mutations on the island of New Guinea
(2020) *PLoS Pathog*, 16 (12), pp. 1-21.
PMID: 33320907
 - Thuy-Nhien, N, Tuyen, NK, Tong, NT, Vy, NT, Thanh, NV, Van, HT
K13 propeller mutations in Plasmodium falciparum populations in regions of malaria endemicity in Vietnam from 2009 to 2016
(2017) *Antimicrob Agents Chemother*, 61 (4).
 - Huang, F, Takala-harrison, S, Jacob, CG, Liu, H, Sun, X, Yang, H
(2015) *MAJOR ARTICLE A Single Mutation in K13 Predominates in Southern China and Is Associated With Delayed Clearance of Plasmodium falciparum Following Artemisinin Treatment*, 212, pp. 1629-1635.
 - Mishra, N, Bharti, RS, Mallick, P, Singh, OP, Srivastava, B, Rana, R
Emerging polymorphisms in falciparum Kelch 13 gene in Northeastern region of India
(2016) *Malar J*, 15 (1), pp. 4-9.
PMID: 27912758
 - Schmedes, SE, Patel, D, Dhal, S, Kelley, J, Szigel, SS, Dimbu, PR
(2021) *Plasmodium falciparum kelch 13*, 27 (7), pp. 2014-2018.
 - Ménard, D, Khim, N, Beghain, J, Adegnik, AA, Shafiul-Alam, M, Amodu, O
A Worldwide Map of Plasmodium falciparum K13-Propeller Polymorphisms
(2016) *New England Journal of Medicine*, 374 (25), pp. 2453-2464.
 - Imwong, M, Hien, TT, Thuy-Nhien, NT, Dondorp, AM, White, NJ.
Spread of a single multidrug resistant malaria parasite lineage (PfPailin) to Vietnam
(2017) *Lancet Infect Dis*, 17 (10), pp. 1022-1023.
[Internet]. PMID: 28948924
 - Witkowski, B, Lelièvre, J, Barragán, MJL, Laurent, V, Su, XZ, Berry, A
Increased tolerance to artemisinin in Plasmodium falciparum is mediated by a quiescence mechanism
(2010) *Antimicrob Agents Chemother*, 54 (5), pp. 1872-1877.
 - Noorazian, Y, Afiqah, SH, Jose Miguel, R, Ridzuan, M, Abd, M, Shuhada, N
(2020) *KELCH13 POLYMORPHISMS IN MALAYSIA (2008–2017)*, 51 (1), p. 2020.
 - Arie, F, Witkowski, B, Amaratunga, C, Beghain, J, Langlois, AC, Khim, N
A molecular marker of artemisinin-resistant Plasmodium falciparum malaria
(2013) *Nature*, 505 (7481), pp. 7481-7485.
[Internet]. 2013 Dec 18 [cited 2023 Oct 12];505 50
 - Coppée, R, Jeffares, DC, Miteva, MA, Sabbagh, A, Clain, J.
Comparative structural and evolutionary analyses predict functional sites in the artemisinin resistance malaria protein K13

- (2019) *Sci Rep*, 9 (1), p. 10675.
[Internet]. PMID: 31337835
- (2018) *Artemisinin resistance and artemisinin-based combination therapy efficacy (Status report—August 2018)*,
[cited 2023 Oct 16]
 - Xie, SC, Ralph, SA, Tilley, L.
K13, the Cytostome, and Artemisinin Resistance
(2020) *Trends Parasitol*, 36 (6), pp. 533-544.
[Internet]. PMID: 32359872
 - Pandit, K, Surolia, N, Bhattacharjee, S, Karmodiya, K.
(2023) *The many paths to artemisinin resistance in Plasmodium falciparum*,
[cited 2023 Oct 19]
 - Tilley, L, Straimer, J, Gnädig, NF, Ralph, SA, Fidock, DA.
Artemisinin Action and Resistance in Plasmodium falciparum
(2016) *Trends Parasitol*, 32 (9), pp. 682-696.
[Internet]
 - Xie, SC, Ralph, SA, Tilley, L.
(2020) *K13, the Cytostome, and Artemisinin Resistance*,
[cited 2023 Oct 11]; PMID: 32359872
 - Yang, T, Yeoh, LM, Tutor, M V., Dixon, MW, McMillan, PJ, Xie, SC
Decreased K13 Abundance Reduces Hemoglobin Catabolism and Proteotoxic Stress, Underpinning Artemisinin Resistance
(2019) *Cell Rep*, 29 (9), pp. 2917-2928.
[Internet]. e5. PMID: 31775055
 - Meshnick, SR, Taylor, TE, Kamchonwongpaisan, S.
Artemisinin and the antimalarial endoperoxides: From herbal remedy to targeted chemotherapy
(1996) *Microbiol Rev*, 60 (2), pp. 301-315.
PMID: 8801435
 - Ivanova, N, Gugleva, V, Dobрева, M, Pehlivanov, I, Stefanov, S, Andonova, V.
We are IntechOpen, the world's leading publisher of Open Access books Built by scientists, for scientists TOP 1%
(2016) *Intech*, i, p. 13.
(tourism)
 - Azman, AN, Fahmirauf, SFM, Fuad, FAA, Hisam, RSR, Yusuf, NAM.
Evaluating the Binding Interactions between Artemisinin and Kelch 13 Protein Mutants Via Molecular Modelling and Docking Studies
(2022) *Journal of Advanced Research in Applied Sciences and Engineering Technology*, 28 (3), pp. 272-286.
 - Fairhurst, RM.
(2016) *Understanding artemisinin-resistant malaria: what a difference a year makes*, 28 (5), pp. 417-425.
 - Suresh, N, Haldar, K.
Mechanisms of artemisinin resistance in Plasmodium falciparum malaria
(2018) *Nature*, 456 (7223), pp. 814-818.
 - Birnbaum, J, Scharf, S, Schmidt, S, Jonscher, E, Maria Hoeijmakers, WA, Flemming, S
A Kelch13-defined endocytosis pathway mediates artemisinin resistance in malaria parasites
(2020) *Science (1979)*, 367 (6473), pp. 51-59.
PMID: 31896710
 - Tsuboi, T, Takeo, S, Sawasaki, T, Torii, M, Endo, Y.
An efficient approach to the production of vaccines against the malaria parasite
(2010) *Methods Mol Biol*, 607, pp. 73-83.
PMID: 20204850

- Müller-Sienerth, N, Shilts, J, Kadir, KA, Yman, V, Homann, MV, Asghar, M
A panel of recombinant proteins from human-infective Plasmodium species for serological surveillance
(2020) *Malar J*, 19 (1), pp. 1-15.
[Internet]
- Birkholtz, LM, Blatch, G, Coetzer, TL, Hoppe, HC, Human, E, Morris, EJ
Heterologous expression of plasmodial proteins for structural studies and functional annotation
(2008) *Malar J*, 7, pp. 1-20.
- Flick, K, Ahuja, S, Chene, A, Bejarano, MT, Chen, Q.
Optimized expression of Plasmodium falciparum erythrocyte membrane protein 1 domains in Escherichia coli
(2004) *Malar J*, 3, pp. 1-8.
- Sony Reddy, K, Pandey, AK, Singh, H, Sahar, T, Emmanuel, A, Chitnis, CE
Bacterially expressed full-length recombinant Plasmodium falciparum RH5 protein binds erythrocytes and elicits potent strain-transcending parasite-neutralizing antibodies
(2014) *Infect Immun*, 82 (1), pp. 152-164.
- Goel, N, Dhiman, K, Kalidas, N, Mukhopadhyay, A, Ashish, F, Bhattacharjee, S.
Plasmodium falciparum Kelch13 and its artemisinin-resistant mutants assemble as hexamers in solution: a SAXS data-driven modelling study
(2022) *FEBS Journal*, 289 (16), pp. 4935-4962.
- Guerra, AP, Calvo, EP, Wasserman, M, Chaparro-Olaya, J.
Production of recombinant proteins from Plasmodium falciparum in Escherichia coli
(2016) *Biomedica*, 36, pp. 97-108.
- Zhao, Y, Lin, YH.
Whole-Cell Protein Identification Using the Concept of Unique Peptides
(2010) *Genomics Proteomics Bioinformatics*, 8 (1), pp. 33-41.
PMID: 20451160
- Siddiqui, FA, Boonhok, R, Cabrera, M, Mbenda, HGN, Wang, M, Min, H
Role of Plasmodium falciparum kelch 13 protein mutations in p. Falciparum populations from northeastern myanmar in mediating artemisinin resistance
(2020) *mBio*, 11 (1), pp. 1-19.
- Abbas, S, Koch, KW.
Quantitative Determination of Ca²⁺-binding to Ca²⁺-sensor Proteins by Isothermal Titration Calorimetry
(2020) *Bio Protoc*, 10 (7), pp. 1-21.
PMID: 33659550
- Pal, B.
Dissection of heme binding to Plasmodium falciparum glyceraldehyde-3-phosphate dehydrogenase using spectroscopic methods and molecular docking
(2017) *Indian J Biochem Biophys*, 54 (1-2), pp. 24-31.
- Kandeel, M, Miyamoto, T, Kitade, Y.
Bioinformatics, enzymologic properties, and comprehensive tracking of Plasmodium falciparum nucleoside diphosphate kinase
(2009) *Biol Pharm Bull*, 32 (8), pp. 1321-1327.
- Boudker, O, Oh, SC.
Isothermal titration calorimetry of ion-coupled membrane transporters
(2015) *Methods*, 76, pp. 171-182.
[Internet]. PMID: 25676707
- Desrosiers, MR, Mittelman, A, Weathers, PJ.
Dried leaf artemisia annua improves bioavailability of artemisinin via cytochrome P450 inhibition and enhances artemisinin efficacy downstream
(2020) *Biomolecules*, 10 (2).
PMID: 32046156

- Bridgford, JL, Xie, SC, Cobbold, SA, Pasaje, CFA, Herrmann, S, Yang, T
Artemisinin kills malaria parasites by damaging proteins and inhibiting the proteasome
(2018) *Nat Commun*, 9 (1), pp. 1-9.
[Internet]

Correspondence Address

Yusuf N.M.; Parasitology Unit, Malaysia; email: noorazian.y@moh.gov.com

Publisher: Public Library of Science

ISSN: 19326203

CODEN: POLNC

PubMed ID: 39146276

Language of Original Document: English

Abbreviated Source Title: PLoS ONE

2-s2.0-85201370409

Document Type: Article

Publication Stage: Final

Source: Scopus

ELSEVIER

Copyright © 2025 Elsevier B.V. All rights reserved. Scopus® is a registered trademark of Elsevier B.V.

 RELX Group™