



ELSEVIER

Contents lists available at ScienceDirect

Data in Brief

journal homepage: www.elsevier.com/locate/dib

Data Article

Preferential binding of 4-hydroxynonenal to lysine residues in specific parasite proteins in plakortin-treated *Plasmodium falciparum*-parasitized red blood cells



Evelin Schwarzer^a, Valentina Gallo^a, Elena Valente^a, Daniela Ulliers^a,
Orazio Tagliatalata-Scafati^b, Paolo Arese^a, Oleksii A. Skorokhod^{*a}

^a Department of Oncology, University of Torino, Via Santena 5bis, 10126 Torino, Italy

^b Department of Pharmacy, University of Napoli "Federico II", Via D. Montesano 49, 80131 Napoli, Italy

ARTICLE INFO

Article history:

Received 13 October 2015

Received in revised form

2 November 2015

Accepted 2 November 2015

Available online 11 November 2015

Keywords:

plakortin

endoperoxide

antimalarial drug

4-hydroxynonenal

post-translational modifications

Plasmodium falciparum

red blood cell

ABSTRACT

The data show the frequencies by which the amino acid residues lysine, histidine and cysteine of six proteins of the malaria parasite *Plasmodium falciparum* are post-translationally modified by the lipoperoxidation endproduct 4-hydroxynonenal after challenging the parasitized red blood cell with plakortin. Plakortin is an antimalarial endoperoxide whose molecular anti-parasitic effect is described in Skorokhod et al. (2015) [1]. Plakortin did not elicit hemoglobin leakage from host red blood cells and did not oxidize reduced glutathione.

© 2015 The Authors. Published by Elsevier Inc. This is an open access article under the CC BY license (<http://creativecommons.org/licenses/by/4.0/>).

Specifications Table

Subject area	Biology
More specific subject area	Molecular pharmacology
Type of data	Tables, graphs

DOI of original article: <http://dx.doi.org/10.1016/j.freeradbiomed.2015.10.399>

* Corresponding author.

E-mail address: olexii.skorokhod@unito.it (O.A. Skorokhod).

<http://dx.doi.org/10.1016/j.dib.2015.11.003>

2352-3409/© 2015 The Authors. Published by Elsevier Inc. This is an open access article under the CC BY license (<http://creativecommons.org/licenses/by/4.0/>).

How data was acquired	MALDI-TOF spectrometer (MALDI micro MX, Waters, Milford, MA, USA); Luminometer Sirius (Berthold, Pforzheim, Germany)
Data format	Processed data
Experimental factors	Parasitized and non-parasitized red blood cells were kept in in vitro cell culture up to 24 h. Chemical extraction for parameter assessment.
Experimental features	Heme-dependent luminol-enhanced luminescence assay for hemoglobin. Thiol-reaction with DTNB and OD measurement at 412 nm. Mass spectrometric analysis of peptides after trypsin-digestion of proteins.
Data source location	University of Torino, Torino, Italy
Data accessibility	Data are provided with this article

Value of the data

- The data show how endoperoxide-elicited oxidative stress might specifically target parasite proteins of functional importance by binding the lipoperoxidation endproduct 4-HNE.
- Data on endoperoxide-elicited modifications of proteins of a unicellular parasite as molecular cause for cell death may be of interest for similar studies with other microorganisms.
- Data for lethal modifications of specific microbial proteins with lipoperoxidation endproducts such as 4-HNE may suggest a novel strategy for high throughput drug research, e.g. antimalarials.
- The data on host red blood cells (RBC) intactness may be useful to monitor host cell damage by redox active substances.

1. Data

1.1. Frequencies of lysine residues modified by 4-HNE

In [Table 1](#) we list the number of lysine, histidine and cysteine residues (K, H and C, respectively) that were found conjugated with 4-HNE and compare them with the total number of each of these amino acids in the respective protein. The complete list of 4-HNE modified *Plasmodium falciparum* proteins detected after plakortin treatment and specific sites of modification are reported in [\[1\]](#).

There is no particular imbalance between the portion of 4-HNE-conjugated amino acids in the proteins extracted from plakortin-treated parasites. We note that *Plasmodium* proteins contain elevated % of asparagine and lysine residues [\[2\]](#), even more than in vertebrates, where K frequency is 2–3 times higher than that of H and C [\[3\]](#).

Evidently, availability of the amino acid residue for 4-HNE is a crucial factor for binding. C residues engaged in disulfide bridges are no ligands for 4-HNE, while the basic amino acid K might be more frequently exposed to the protein–solvent interface compared to H, and hence be more accessible to 4-HNE binding [\[4\]](#). The data about unbalanced ratios were published for other proteins such as liver fatty acid-binding protein [\[5\]](#), cytochrome c [\[6\]](#) and alpha-synuclein [\[7\]](#), where 4-HNE binding site distribution between K, H, C was 4:1:1 [\[5\]](#); 10:1:0 [\[6\]](#) and 2:1:0 [\[7\]](#) respectively.

1.2. Data for hemoglobin release from RBC under plakortin treatment

Culture of RBC infected (parasitized) with trophozoite-stage *P. falciparum* were treated with 0–10 μM of plakortin and hemoglobin release was measured in culture supernatant ([Fig. 1](#)). The concentration is indicated in nmol/l. The concentration in the whole RBC suspension was 1 mM and assessed after complete lysis of RBC in NaOH/Triton: this is the highest maximal achievable concentration of hemoglobin in the supernatant corresponding to 100% RBC lysis.

As shown in [Fig. 1](#) the hemoglobin was not released from RBC even at 10 μM plakortin, the highest applied concentration. Very low lysis was detectable in non-parasitized RBC (npRBC) and a still very modest, although double as high value in trophozoite-parasitized mature forms of *P. falciparum*. This

Download English Version:

<https://daneshyari.com/en/article/174931>

Download Persian Version:

<https://daneshyari.com/article/174931>

[Daneshyari.com](https://daneshyari.com)