

Detection of histidine rich protein & lactate dehydrogenase of *Plasmodium falciparum* in malaria patients by sandwich ELISA using in-house reagents

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Background & objectives: Despite major control efforts, malaria remains a major public health problem that still causes high mortality rate worldwide especially in Africa and Asia. Accurate and confirmatory diagnosis before treatment initiation is the only way to control the disease. The present study was undertaken to develop reagents using sandwich ELISA for simultaneous detection of PfHRP2 (*Plasmodium falciparum* histidine rich protein) and PflDH (*P. falciparum* lactate dehydrogenase) antigens in the proven malaria cases.

Methods: The antibodies were raised against two epitopes of PfHRP2 protein and three unique and unexplored epitopes of PflDH protein. These antibodies were able to detect PfHRP2 and PflDH antigens in culture supernatant and parasitized RBC lysate of *P. falciparum*, respectively up to 50 parasites/ μ l. The in-house reagents were tested in 200 *P. falciparum* positive patients residing in Baghpat district of Uttar Pradesh in northern India.

Results: Microsphere (PLGA) with CpG ODN were used to generate high titre and high affinity antibodies against selected peptides of PfHRP-2 and pLDH antigen in mice and rabbit. The peptide specific peak titre varied from 12,800 - 102,400 with an affinity ranging 0.73 - 3.0 mM. The indigenously developed reagents are able to detect PfHRP2 and PflDH antigens as low as 75 parasites/ μ l of blood with a very high sensitivity (96-100%) and specificity (100%).

Interpretation & conclusions: The study highlight the identification of unique epitopes of PfHRP2 and PflDH, and the generated antibodies against these antigens were used for quantitative estimation of these two antigens using sandwich ELISA. No crossreactivity with *P. vivax* infected patients was observed with the sera.

Key words Diagnosis - ELISA - histidine rich protein 2 - lactate dehydrogenase - malaria - *Plasmodium falciparum* (Pf)

Malaria is potentially a life-threatening disease, which predominantly occurs in tropical and subtropical regions. It continues to be a major public health problem in endemic countries lacking adequate health care and malaria control programme.

The clinical diagnosis based on symptoms is the most common method used but it shows less specificity due to overlapping of symptoms with other tropical diseases. Microscopic examination of stained blood films is another cornerstone for diagnosis and estimation of parasite load in malaria victims. Although it is cost effective and simple, but it shows poor reproducibility, variable sensitivity and requires skilled operators for accurate diagnosis. Importantly, in remote areas where malaria commonly occurs, it is hard to maintain good quality microscopy and the result based on microscopic examination is sometimes unreliable. In many cases co-infection with *Plasmodium falciparum* and *P. vivax* occur, therefore, accurate and prompt parasitological confirmation of malaria infection is essential for effective disease management.

Many of the new technologies for malaria diagnosis incorporate immunochromatographic procedure, where conjugated monoclonal antibodies are the key reagents. Currently many rapid diagnostic tests (RDTs) are widely used for the diagnosis of malaria. These RDTs are simple lateral-flow immunochromatographic tests that detect parasite specific antigens released from red blood cells. Two of the tests, the ICT Malaria Pf/Pv and ParaSight-F detect histidine rich protein 2 (HRP-2), a protein produced by asexual stages and young gametocyte of *P. falciparum*^{1,2}. The third test OptiMAL detects *Plasmodium* lactate dehydrogenase (pLDH), a marker protein for the intraerythrocytic form of the malaria parasite. HRP-2 is an abundant protein produced by all blood stages of *P. falciparum*³. However, there are certain limitations of these rapid tests, including the decrease in sensitivity⁴ (70 in parasitaemia <50/μl) and also these tests cannot give quantitative (parasite/μl) results with malaria positive serum samples.

The HRP-2 antigen is expressed by all *P. falciparum* parasites regardless of knob-phenotype, and can be recovered from culture supernatants as a secreted soluble protein⁵. HRP-2 can be detected in erythrocytes, serum, plasma, cerebro-spinal fluid and even in urine^{6,7}. Sequencing of the genomic DNA has shown that HRP-2 antigen contains 35 per cent histidine as well as alanine and aspartate (40 and 12%, respectively).

It is characterized by the presence of tandem repeats of AHH and AHHAAD. Since HRP-2 antigen is only produced by *P. falciparum*, these tests cannot be used for the detection of *P. vivax* or other human malarial parasites. In endemic areas *P. falciparum* has been reported to lack HRP-2 or HRP-3 or both in positive patients⁸. In India, *P. falciparum* lacking PfHRP-2 and the PfHRP-3 gene has so far not been reported. However, the HRP-2 antigen remains detectable for several weeks after parasite clearance which causes false-positive RDT results in the test samples⁹.

Distribution of *P. falciparum* isolates lacking PfHRP-2 and PfHRP-3 is a major challenge for RDTs. In these areas parasite specific pLDH is the major target for the detection of malaria. pLDH is an intracellular glycolytic enzyme, which catalyses the oxidation of lactate to pyruvate. In *P. falciparum*, the coenzyme for LDH is preferably 3-acetyl pyridine adenine dinucleotide, (APAD), whereas the activity of human LDH requires β nicotinamide adenine dinucleotide¹⁰ indicating that pLDH could be a good marker following active malarial infections¹¹. pLDH is expressed at high levels during asexual stage or blood-stage in all four malaria parasites¹² and also correlated with the number of parasites present in the plasma of infected patients¹³. pLDH from *P. vivax*, *P. malariae*, and *P. ovale* exhibit 87 per cent sequence identity with pLDH from *P. falciparum*¹⁴. Genetic diversity may be particularly important for PfHRP-2-based RDTs since the antigen consists of a number of alanine and histidine-rich amino acid repeats and varies in size between parasite strains¹⁵.

In the present study we developed a simple antigen capture ELISA for detection of *P. falciparum* specific HRP-2 and LDH in the blood of malaria patients using in-house reagents.

Material & Methods

The study was conducted in the Department of Biochemistry, All India Institute of Medical Sciences (AIIMS), New Delhi, India, and the study protocol was approved by the Ethics Committee on animal experimentation, AIIMS.

Identification and synthesis of peptides and adjuvant: The protein sequence for PfLDH (PBD id PF3D7_1324900) and PvLDH (PVX_116630) were obtained from the PlasmoDB version 9.2 malaria database (www.plasmodb.org). DNASTar and Bcelpred software (www.IMTECHBioinformatics) were used for

the selection of peptides on the basis of physicochemical properties such as antigenicity, hydrophilicity, hydrophobicity, flexibility/mobility, exposed surface antigenicity and β turns. Structural data for epitope location were drawn by PYMOL [DeLano WL (2002) The PYMOL Molecular Graphics System on world wide web <http://www.pymol.org>]. Alignments were performed by EMBOSS Stretcher protein alignment EMBL-EBI software (www.ebi.ac.uk).

Two peptides with tandem repeat sequence from the *Pf*HRP-2 protein, and three unique peptide sequences from LDH protein were selected. The sequences were AHH (AHHAAD)₄ (HRP-2 peptide I), (AHHA)₄ (HRP-2 peptide II) and (LFDIVKNMPHGKALDTSHT)₂ (LDH peptide I), (TNVMAYSNCKVSGSNTYDDLA)₂ (LDH peptide II), (VVLGANGVEQVIELQLNSEEKA)₂ (LDH peptide III).

All peptides were synthesized by solid phase peptide synthesis using F-moc chemistry¹⁶. After analyzing all the above peptide sequences of *Pf* LDH by BLAST (<http://blast.ncbi.nlm.nih.gov>), no sequence similarity with the other vertebrate LDH including humans was observed. The purity of each peptide was assessed by amino acid analysis and HPLC and after purification these were found to be >95 per cent pure. The CpG oligodeoxynucleotide (ODN) 1826 of B class (also known as K type) with a nuclease resistant phosphorothioate backbone was used as an adjuvant to enhance antibody response. It was procured commercially from Coley pharmaceuticals (Wellesley, USA).

Preparation of CpG ODN microparticles (Microspheres): Poly (poly-DL-lactide-co-glycolide) PLGA microparticles entrapping peptide antigens with CpG ODN [peptide: CpG, (6:1) w/w] were prepared using water in oil-in water emulsion and solvent evaporation method¹⁷. The percentage entrapment for different peptides was in the range of 50-60 per cent as determined by BCA (bicinchoninic protein assay)¹⁸. The size distribution of microsphere was determined by laser diffraction (Malvern Instrument, UK) and the size was between 5-10 μ m. Microsphere morphology was also studied by scanning electron microscopy (Philips, CM 10, USA). The percentage entrapment of CpG ODN was determined by extracting CpG ODN from microparticle in 10mM TE (Tris-EDTA) buffer, pH 8.3 as per protocol¹⁹.

Animals, immunization and serum collection: Six to eight week old inbred BALB/c (H-2^d) mice and three months old rabbit were procured from the Experimental Animal Facility, AIIMS, New Delhi, India. A group of six mice were immunized subcutaneously at the foot-pad with *Pf*HRP-2 peptide I and *Pf*LDH peptides [regions I (33-51aa), II (51-71 aa) and III (280-306aa)], respectively in microsphere containing CpG ODN (5 μ g) with primary dose of 30 μ g on day 0 followed by a booster dose of 20 μ g with 2.5 μ g CpG ODN on days 32 and 45. Likewise, four rabbits were also immunized subcutaneously at multiple site with *Pf*HRP-2 peptide II with primary dose of 100 μ g with 20 μ g CpG ODN on day 0 followed by a booster dose of 50 μ g with 10 μ g CpG ODN on days 32 and 45. The animals were provided with food and water *ad libitum*. Serum was collected on days 15, 28, 42, 60 and 90.

ELISA: The peptide specific antibody and peak titres were estimated using standard ELISA protocol. Briefly, 100 ng/100 μ l of the peptide was coated per well and incubated with serum sample (mice/rabbit serum) at a standardized dilution (1:100 v/v) for serum IgG levels as well as two-fold serial dilution for measurement of peak antibody levels as per our reported protocol²⁰. The end point titres were expressed as the reciprocal of the highest serum dilution giving an absorbance = pre-immune serum +4SD.

Purification and biotinylation of antibodies raised against synthetic peptides of PfHRP-2 and PfLDH protein: Initially albumin was removed from serum by ammonium sulphate (40-50% sodium) precipitation or "salting out" procedure²¹ and subsequently, Sepharose Protein A column was used to further purify the enriched IgG fraction. The pooled fractions were dialyzed against 0.01 M PBS (pH 7.4). The dialysate was concentrated and the IgG amount was determined using the BCA method¹⁸.

Biotinylation of the purified antibodies was done using N-hydroxysuccinimide ester of biotin (Sigma Immuno Probe Biotinylation reagents, USA) according to the manufacture's protocol Further, biotinylated antibodies were purified by gel filtration procedure (G-25 column). The extent of biotinylation and the ratio of biotin to antibody was determined by the avidin-HABA (4'-Hydroxazobenzene-2-carboxylic-acid) assay²², 2-3 biotin molecules were found to be linked per molecules of IgG.

Binding affinity of the anti-peptide antibodies (K_D): The affinity of antibodies raised against different peptides

in mice was determined by measuring the dissociation constant (K_D)²³. In brief, immunized serum from mice/rabbit at 1 : 200 dilutions was incubated with different concentrations of the peptide (0.1-10 nM) for 15 h at 20°C so as to attain antigen-antibody equilibrium. The antigen-antibody complexes were transferred onto the wells of the microtitre plates previously coated with the respective peptide (500 ng/well). The plates were incubated for 90 min at 37°C. After washing 3 times with PBS-T, goat anti-mouse/rabbit IgG HRPO (horse radish peroxidase) conjugate was (1:1000) added (100 µl/well) and incubated for 1 h at 37°C. Dissociation constants were calculated using regression analysis and a simplification of the mathematical equation of Scatchard and Klotz²⁴.

$$\frac{A_0}{A_0 - A} = 1 + \frac{K_D}{a_0}$$

Where A_0 is the absorbance without free antigen, A is the absorbance with free antigen and a_0 is the total amount of antigen added to the reaction mixture.

Cultivation of P. falciparum and preparation of P. falciparum lysate: *P. falciparum* isolates (Indian isolates FDL-B, FDL-NG, FSH-4 and FSH-11) were maintained in *in vitro* cultures using O⁺ RBCs and AB⁺ serum²⁵. Antigen was prepared from cultures enriched with late trophozoite and schizonts. Parasites were freed by saponin lysis and soluble extract was obtained after sonication at 14 µA for 90 sec. Each batch of culture was monitored for parasitaemia by microscopy, then parasites were harvested and culture supernatants were aspirated. Parasitized RBCs were washed thrice with PBS, and both the pellet and supernatant were stored at -20°C.

Study area and study population: A cross-sectional survey was conducted in the villages of the Baghpat district of Uttar Pradesh (UP), in northern India, from August 2010 to November 2011. This area is endemic for malaria having seasonal transmission of both *P. vivax* and *P. falciparum*. Early and prolonged monsoons are responsible for intensive transmission for both the species. All the patients were recruited after taking written informed consent from them. Overall 2050 individuals were screened for malaria infection. Giemsa stained thick and thin peripheral blood smears of each patient were examined by trained microscopist. The malaria positive samples were provided to us by Dr Sukla Biswas, NIMR, New Delhi after taking the ethical clearance from NIMR, New Delhi. All these samples were collected with the help of Doctor on duty of the primary health centre,

Bhagpat, U.P. In addition, finger-prick blood samples of malaria positive and healthy subjects were tested by immunochromatography based rapid diagnostic test (RDT) kit, FalciVax, rapid test for *Pv/Pf* (Zephyr Biomedicals, Verna, Goa, India). Parasite density was estimated by counting the number of parasites per 200 leukocytes and the counts were converted to number of parasites/µl blood taking 8000 leukocytes/µl as a standard mean. More than 100 microscopic thick smear fields were checked before declaring a slide negative.

Two hundred confirmed malaria patients from the above samples were included in the study. Blood samples from patients with uncomplicated *P. falciparum* infection were collected by finger-prick in heparinized tubes. In a similar way, blood samples were also collected from 50 cases of *P. vivax* infected malaria and 50 healthy individuals to serve as a negative control. Patients diagnosed with malaria were treated with recommended antimalarials as per National Malaria Eradication Programme. All the samples were transported at 4°C to NIMR, New Delhi.

Development of PfHRP2 and PfLDH antigen assay design: For detecting the PfHRP2 antigen levels in *P. falciparum* positive patients, an ELISA was designed in which the plates were coated overnight at 4°C with 2.5 µg/100 µl of purified anti-PfHRP2 peptide I antibody raised in mice. After blocking with 5 per cent BSA (bovine serum albumin), 100 µl of RBC lysates of the blood samples from *P. falciparum* positive patients were added in 1:100 dilutions in each well and incubated at 37°C for 1 h. Plates were washed with PBS-Tween-20 and then 2.5 µg/100µl of purified biotinylated rabbit anti-PfHRP2 peptide II antibody was added and incubated at 37°C for 1h. For detecting parasite LDH antigen levels in the same patients, an ELISA was designed in which plates were coated with 2.5 µg of purified anti-PfLDH peptide III antibody in coating buffer and kept overnight. After blocking with 5 per cent BSA, 100 µl of RBC lysates of the blood samples from *P. falciparum* positive patients were added in 1:100 dilution in each well and incubated at 37°C for 1 h. Plates were washed with PBS-Tween-20 and then a cocktail of 2.5 µg of purified biotinylated anti-PfLDH peptide I and anti-PfLDH peptide II antibody was added and incubated at 37°C for 1 h. The bound antigen-antibody complex was detected using streptavidin-HRPO (1:1000 dilutions). Colour was developed using ortho pherylene diamine (OPD) as a chromogen and the absorbance was read at 492 nm. Similar ELISA was done with *Pf* culture supernatant at different dilutions

(neat, 1:2, 1:4, 1:8, 1:16, 1:32 and 1:64) or parasitized RBC lysate at various parasitaemia levels (ranging from 250,000-50 parasites/ μ l) to develop a standard curve. RBC lysate and culture supernatant from non-infected blood samples of normal volunteers were used as control.

Statistical analysis: The data analysis was done with the help of Stata/IC (version 12.1) Stata Corporation College, Stations, Texas, USA. ELISA result was analyzed by Rank-Sum test (non-parametric approach) to compare the values of HRP-2 and LDH antigen between cases and control and comparison between two sets of ELISA and microscopy was done by Spearman Rank correlation coefficient. Antibody titres of different peptides from different bleeds were determined by the Friedman test separately for each group and group variability was analyzed by Kruskal-Wallis test (data not shown). The slope of the lines (K_D values) was calculated by regression analysis.

Results

Selection of peptides for generating high titre antibody in animal models: Three regions from *p*LDH were identified that showed amino acid residues differed between *P. falciparum* and *P. vivax* (Fig. 1). All three selected regions showing four to six amino acid difference were thought to be sufficient to generate specific antibodies against each selected LDH peptide and also able to differentiate one malaria LDH from another malaria species.

Surface localization of the three selected peptides was necessary to evaluate the accessibility of raising anti-peptide antibody for native antigen (*p*LDH protein). Surface localization of the peptides was by a PYMOL software program (Fig. 2) illustrated that all peptides were found to be located on the surface of the native protein.

Serum anti-peptide humoral response in microparticle formulation of P. falciparum LDH and HRP-2 antigens: The antibody responses to all the five peptides raised in mice and rabbits were measured as a proportion of peak antibody titres. Peptides entrapped in PLGA microspheres with the CpG ODN generated peptide specific high antibody levels in all the five bleeds and the levels were maintained till 90 days post-immunization. IgG peak titre for *Pf*HRP-2 peptide I ranged 51,200 - 1, 02,400 on days 28 and 42 (Table I). Mice/rabbits immunized with *Pf*LDH peptide I and *Pf*HRP-2 peptide II also showed peak antibody levels 51,200 on days 28 and 42 and the titres fell to 25,600 on day 90. Similarly for *Pf*LDH peptide II antisera, IgG peak titres were in the range of 51,200 - 1,02,400 on days 28 and 42, then gradually declined by day 90 to 25,600. For *Pf*LDH peptide III antisera, IgG peak titres were in the range 25,600 - 51,200 on days 28 and 42 which persisted up to day 90 (Table I).

*Dissociation constant (K_D) of *Pf*HRP-2 peptide antisera and *Pf*LDH peptides antisera:* The K_D value of anti-*Pf*HRP-2 peptide I antisera (0.73 nM) was lower than

<i>Pf</i> LDH	MAPKAKIVLVGSGMIGGVMATLIVQKNLGDVV LFDIVKNMPHGKALDTSH	50
<i>Pv</i> LDH	MTPKPKIVLVGSGMIGGVMATLIVQKNLGDVVMFDVVKNMPQGKALDTSH	50
<i>Pf</i> LDH	TNVMAYSNCKVSGSNTYDDL AGADVVIIVTAGFTKAPGKSDKEWNRDILLP	100
<i>Pv</i> LDH	SNVMAYSNCKVTGSNSYDDLKGADVVIIVTAGFTKAPGKSDKEWNRDILLP	100
<i>Pf</i> LDH	LNNKIMIEIGGHIKKNCPNAFIIVVTNPVDVMVQLLHQHSGVPRNKIIGL	150
<i>Pv</i> LDH	LNNKIMIEIGGHIKKNCPNAFIIVVTNPVDVMVQLLFEHSGVPRNKIIGL	150
<i>Pf</i> LDH	GGVLDTSRLKYYISQKLNVCPRDVNAHIVGAHGNKMVLLKRYITVGGIPL	200
<i>Pv</i> LDH	GGVLDTSRLKYYISQKLNVCPRDVNALIVGAHGNKMVLLKRYITVGGIPL	200
<i>Pf</i> LDH	QEFINNKLISDAELEAIFDRTVNTALEIVNLHASPYVAPAAAIIEMAESY	250
<i>Pv</i> LDH	QEFINNKITDEEVEGIFDRTVNTALEIVNLLASPYVAPAAAIIEMAESY	250
<i>Pf</i> LDH	LKDLKQVLCSTLLEGQYGHSDIFGGT PVVLGANGVEQVIELQLNSEEKA	300
<i>Pv</i> LDH	LKDIKQVLCSTLLEGQYGHSDIFGGT PLVIGGTGVEQVIELQLNAEET	300
<i>Pf</i> LDH	KFDEAIAETKRMKALA	316
<i>Pv</i> LDH	KFDEAVAETKRMKALI	316

Fig. 1. EMBOSS stretcher alignment of *Plasmodium* lactate dehydrogenase amino acid sequences. The unique plasmodial epitope differentiating *Pf*LDH and *Pv*LDH appear below the highlighted region. Accession numbers are *Pf*LDH: PlasmoDBid: PF3D7_1324900), *Pv*LDH: PlasmoDBid: PVX_116630.

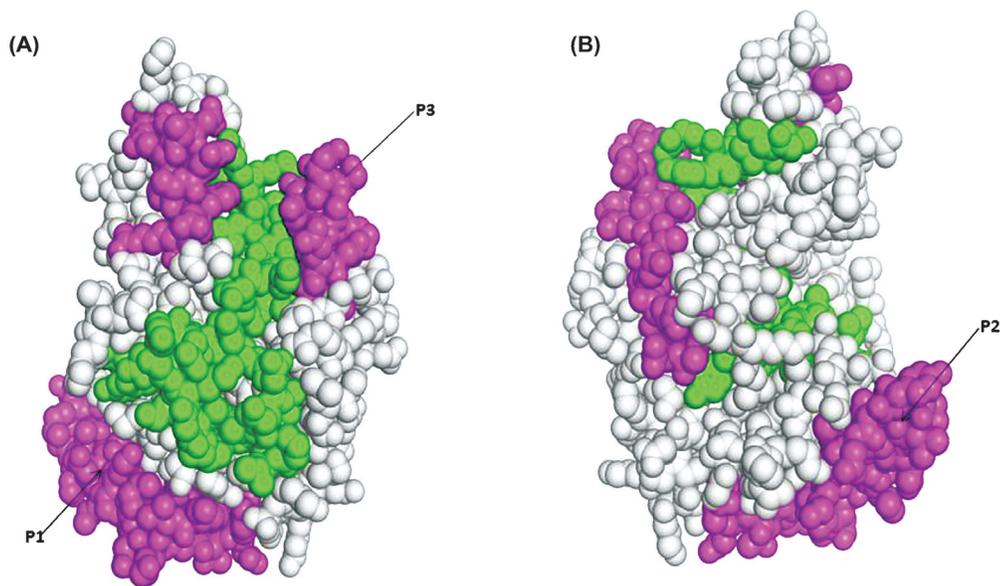


Fig. 2. Schematic diagram showing the localization of *P. falciparum* specific LDH epitope in the pLDH protein at (A) 0° rotation and (B) 180° rotation along the Y axis (PDB Id: 2A94). The peptide localization of all *Pf*LDH peptide was indicated in magenta. The Figure was drawn by PyMOL [DeLano WL (2002) The PyMOL molecular graphic system on world wide Web: <http://www.pymol.org>, for this study.

anti-*Pf*HRP-2 peptide II antisera (1.5 nM) and the K_D value of anti-*Pf*LDH peptide II antisera (1.2 nM) was also lower as compared to the anti-*Pf*LDH peptide I and III antisera (1.4 and 3.0 nM). Thus all the peptides generated high titre and high affinity antibodies.

Quantification of *Pf*HRP-2 antigen in *P. falciparum* culture supernatant or parasitized RBC lysates using *Pf*HRP-2-based ELISA: The *Pf*HRP2 assay allowed the detection of HRP-2 antigen in the culture supernatant at dilution (neat, 1:2, 1:4, 1:8, 1:16, 1:32 and 1:64) of all the four isolates of *P. falciparum* (Fig. 3) and the parasitized RBC lysate (250,000-50 parasites/ μ l) (Fig. 4A). The lower limit of *Pf*HRP-2 antigen detection (50 parasites/ μ l) in parasitized RBC lysate was better than the detection limit by light microscopy which gave 100 per cent sensitivity

when the number of parasites/ μ l of blood was >50. The number of parasites/ μ l blood was found to be positively correlated with *Pf*LDH antigenaemia ($r=0.863$, $P<0.001$) and in the infected RBC (IRBC) lysates ($r=0.839$, $P<0.001$). The assay was repeated thrice to confirm reproducibility.

Quantification of LDH antigen in *P. falciparum* culture supernatant or IRBC lysates using *Pf*LDH-based ELISA: The *Pf*LDH assay allowed the detection of *Pf* LDH antigen in the parasitized RBC lysate at parasitaemia levels ranging 250,000-50 parasites/ μ l (Fig. 4B). The lower limit of *Pf*LDH antigen detection (50 parasites/ μ l) in parasitized RBC lysate was better than the detection limit by light microscopy which gave 100 per cent sensitivity when the number of parasites/ μ l of blood was ≥ 50 . The number of parasites/ μ l blood

Table I. End point titres (in thousands) of antibodies raised against the peptides of *Pf*HRP2 and *Pf*LDH antigens

	Day 15	Day 28	Day 42	Day 60	Day 90
<i>Pf</i> HRP2 Peptide I	25,600	51,200	1,02,400	1,02,400	51,200
<i>Pf</i> HRP2 Peptide II	25,600	51,200	1,02,400	1,02,400	1,02,400
<i>Pf</i> LDH Peptide I	25,600	51,200	51,200	25,600	25,600
<i>Pf</i> LDH Peptide II	25,600	51,200	1,02,400	1,02,400	25,600
<i>Pf</i> LDH Peptide III	12,800	25,600	51,200	51,200	25,600

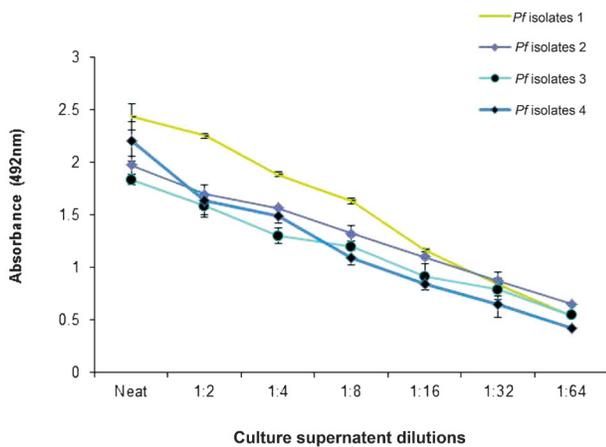


Fig. 3. Quantification of *Pf*HRP2 antigen in culture supernatants of four different isolates of *P. falciparum* (Isolate 1 FDL-B, Isolate 2 FDL-NG, Isolate 3 FSH-4 and Isolate 4 FSH-11). Values are mean + SD of 3 observations.

were found to be positively correlated with *Pf*LDH antigenaemia ($r=0.831, P<0.001$) in the infected RBC lysates. We did not perceive any detectable amount of pLDH in culture supernatants. The assay was repeated thrice for reproducibility.

Detection of Pf HRP-2 antigen in blood of malaria patients by PfHRP-2 assay: Two hundred malaria positive serum samples confirmed by microscopy and RDT were studied to evaluate the newly developed antigen capture assay for *Pf*HRP-2 and *Pf*LDH. The results are presented in Table II. In this study, at 95% CI the specificity of *Pf*HRP-2 assays for

diagnosis of *P. falciparum* parasites were 100 per cent (91.1-100%) and sensitivity was in the range 96 per cent (77.7-99.8%) to 100 per cent (96.4-100%) when the parasitaemia was 0.0015 to 0.015 per cent, respectively. The per cent positive predictive values (PPV%) were found to be 74.7 to 100 per cent, the negative predictive values % (NPV%) were found to be 91.1 to 100 per cent and area under curve (AUC) was found to be 1.00 when parasite/ μ l >10,000-150. However, PPV% was calculated to be 82.8-100 per cent, NPV% was 98.04 per cent (88.2-99.2%) when parasite/ μ l >75-150 and AUC was found to be 0.98. This assay was able to detect the *Pf*HRP-2 antigen in the blood of all the subjects with parasitaemia level ranging 75 to 250, 000 parasites/ μ l blood which corresponds to 0.0015 to 5 per cent parasitaemia (Fig. 5A). When the association between number of parasites/ μ l blood and HRP-2 antigenemia was measured, the level of parasitaemia were found to be positively correlated with *Pf*HRP2 antigenemia ($r=0.862, P<0.001$). When all 200 patients of proven malaria infection were divided into different groups based on parasite/ μ l blood, the association between mean parasitemia levels in different groups and the corresponding mean *Pf*HRP-2 antigenemia was also positively correlated ($r=0.979, P<0.001$). As the parasitemia level increased, the *Pf*HRP-2 concentration was found to be increased (Fig. 5A, 5B). The indigenously developed antibodies were able to detect *Pf*HRP-2 in these blood samples at parasitaemia of approximately 75 parasites/ μ l (Table II). Fifty of *P. vivax* positive cases and 50 healthy, smear negative individuals served as control and had undetectable

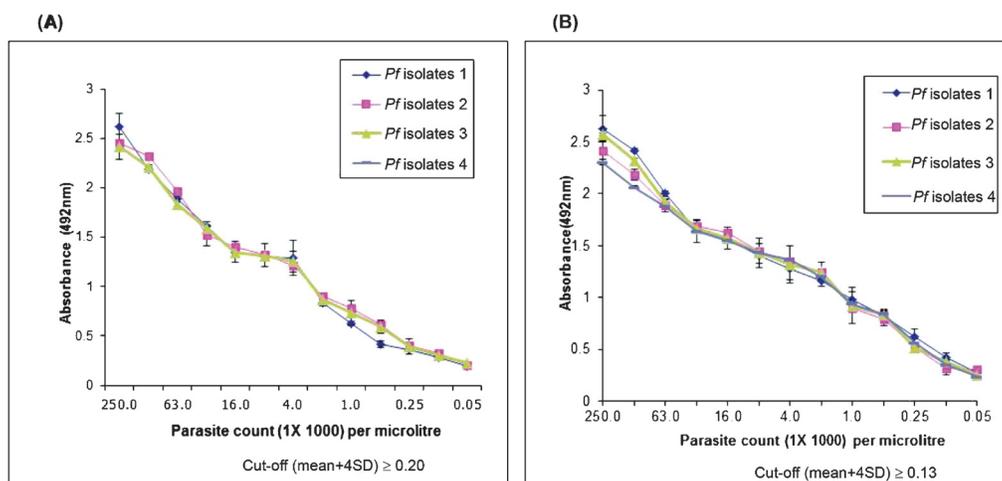


Fig. 4. Quantification of *Pf*HRP2 antigen (A) and *Pf*LDH antigen (B) in parasitized RBC lysates of four different isolates of *P. falciparum* (Isolate 1 FDL-B, Isolate 2 FDL-NG, Isolate 3 FSH-4 and Isolate 4 FSH-11). Values are mean + SD of 3 observations.

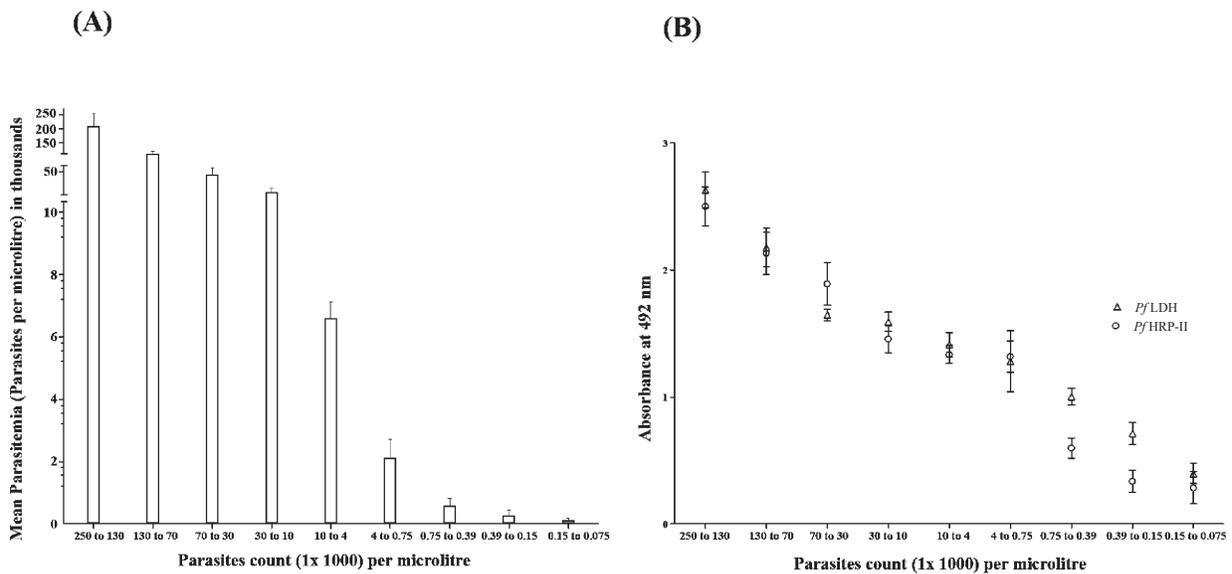


Fig. 5. (A) Association between mean parasitaemia and parasite count in different groups of *P. falciparum* positive patients. Parasitaemia is expressed as parasites/ μ l and data expressed as mean +4 SD. (B) Relationship between parasite count and mean *Pf*HRP2 (O) and mean *Pf*LDH antigen concentration (Δ) in *P. falciparum* positive. Parasitemia is expressed as parasites/ μ l and data is expressed as mean + 4 SD.

levels of *Pf*HRP antigen. Cut-off for positivity was calculated as \geq mean + 4SD of negative control. A sample is given an absorbance (Cut-off) \geq 0.20 was considered positive for *Pf*HRP-2 antigen.

Detection of Pf LDH antigen in blood of malaria patients by Pf LDH assay: Blood samples from 200 malaria patients were also examined for *Pf*LDH and compared with microscopy and RDT. The specificity of *Pf*LDH assays for diagnosis of *P. falciparum* parasites was 100 per cent (91.1-100%) and sensitivity was 100 per cent (83.4-100%) at 95% CI. The PPV% values were found to be 83.4-100 per cent, the NPV% values were found to be 91.1 to 100 per cent and AUC was 1.00 when parasite/ μ l $>$ 75 which corresponded

to $>$ 0.0015 per cent parasitaemia. The developed in-house assay detected *Pf*LDH antigen in the blood of all the subjects with parasitaemia level ranging from 75 to 250,000 parasites/ μ l blood, which corresponded to 0.0015 to 5 per cent parasitaemia (Fig. 5A). When the association between number of parasites/ μ l and *Pf*LDH antigenaemia was measured, the level of parasitaemia were found to be positively correlated with *Pf*LDH antigenaemia ($r=0.878$, $P<0.001$) (Fig. 5B). The associations between mean parasitaemia levels in different groups of patients as described above and the corresponding mean *Pf*LDH antigenaemia were also found positively correlated ($r=0.972$, $P<0.001$). As the parasitaemia level increased, the *Pf*LDH concentration in the blood was also found to be increased in these

Table II. Performance of *Pf*HRP2- and *Pf*LDH- based ELISA on *P. falciparum* confirmed serum samples of patients from Baghpat district of Uttar Pradesh

Blood film % parasitaemia	Parasites/ μ l	Total	<i>Pf</i> HRP2 assay	Sensitivity (%)	<i>Pf</i> LDH assay	Sensitivity (%)	Specificity (%)
$>$ 0.2	$>$ 10,000	128	128	100	128	100	100
0.2-0.015	10,000-750	17	17	100	17	100	100
0.015-0.008	750-390	15	15	100	15	100	100
$>$ 0.003	390-150	15	15	100	15	100	100
$>$ 0.0015	150-75	25	24	96	25	100	100
<i>P. vivax</i> $>$ 0.015	0	50	0	0	0	0	-
Negative	0	50	0	0	0	0	-

patients (Fig. 5B). Thus, the indigenously developed antibodies detected *Pf*LDH in these patient samples at parasitaemia of approximately 75 parasites/ μ l of blood. A sample is given an absorbance (Cut-off) ≥ 0.13 was considered positive for *Pf*LDH antigen detection. The developed reagents were also tested with 50 samples of *P. vivax* positive serum. It did not show any reactivity with the *P. vivax* samples. The assay showed undetectable levels of *Pf*LDH antigen in 50 healthy and smear negative controls.

Discussion

In the present study, the performance of antibodies raised against two epitopes for *Pf*HRP-2 and three different epitopes for *Pf*LDH antigens was assessed for their use as immunoreagents for development of ELISA-based diagnostic assay to assess the antigen load in a large number of *Pf* infected serum samples. All the selected regions were surface exposed to the native protein and hence generated antibodies against these peptides and thus was able to differentiate between *P. falciparum* and *P. vivax*.

The use of the core characteristics of *Plasmodium* HRP-2 and LDH antigens for detection of malaria, for estimation of parasite biomass^{5,26}, in RDT²⁷ and in antigen-antibody assay has been reported²⁸. This feature was exploited in the designing of first immunodiagnostic assay for *P. falciparum*-specific HRP-2²⁹ and later the second generation immunochromatographic based rapid diagnostic tests³⁰ were developed for *P. falciparum*. Studies on sequence variability among *Pf*LDH isoforms from different strains showed low diversity, which suggested that the antigen conserved its sequence³¹. The characteristics of the core technology for *Pf*LDH have been outlined elsewhere¹³; these studies used dipstick and ELISA to evaluate the characteristic of *Pf*LDH. It was shown that *Pf*LDH dipstick had a threshold effect of approximately 200 parasites/ μ l of blood. Levels of *Pf*LDH were more consistent with peripheral parasitaemia than *Pf*HRP-2.

Thus, we have selected peptides from two regions of HRP-2 and three unique and unexplored region of pLDH based on the structural, peak antibody titres and affinity values for different antibodies, the antisera of *Pf*HRP-2 peptide I and *Pf*LDH peptide III (278-300 amino acid regions) were used as capture antibody and the antisera of *Pf*HRP-2 peptide II and a cocktail mixture of *Pf*LDH peptide I (33-51 amino acid region) and *Pf*LDH peptide II (51-71 amino acid regions) were used as detecting antibody in the enzyme

immunoassay. Also to enhance the immunogenicity, the microencapsulated peptides were co-entrapped with CpG oligodeoxynucleotide as an adjuvant. It is well known that CpG adjuvant is more potent when used in conjunction with the delivery system largely as a consequence of its improved delivery into the endosome of antigen presenting cell (APC), where it can more easily interact with TLR9 (Toll like receptor 9)³². Hence in this study, it was observed that when the length of peptide sequences of *Pf*HRP-2 and *Pf*LDH proteins was increased by tandem repeat and CpG was combined with the microsphere delivery, the dissociation constant of antisera was considerably reduced thereby indicating that antibodies have high affinity, which helped to detect low parasitaemia levels up to ~ 75 parasites/ μ l blood in *P. falciparum* positive serum. The antibodies developed against *Plasmodium* lactate dehydrogenase were used in the differential diagnosis of *P. falciparum* and *P. vivax* infection. The lower limits of detection for *Pf*HRP-2 and *Pf*LDH were comparable to those reported for other rapid diagnostic tests³³. Also, *Pf*LDH and *Pf*HRP-2 levels detected in the present study can be used to complement microscopy in the target population. Microscopy only detects circulating infected RBC and is not an accurate measure of parasite burden, particularly in *P. falciparum* infections where mature stages are known to be sequestered in various tissues. It is now widely accepted that *Pf*LDH levels reflect current infection whereas *Pf*HRP-2 levels indicate both past and current infection⁹. Thus concurrent measurement of these two antigens by ELISA provides a better evaluation of parasite burden especially in areas, where malaria is endemic³⁰. There are a few RDT kits commercially available for the diagnosis of *P. falciparum* specific HRP-2 and LDH antigen²⁷. The overall sensitivity reported was 88.7 per cent with blood parasite count < 100 parasite/ μ l and increased from 94.3 to 99.3 per cent when the blood parasite count was > 100 -10000 parasite/ μ l with a specificity of 97.5 per cent³⁴. However, in the present study certain unique epitopes of *Pf*HRP-2 and *Pf*LDH were identified and used for diagnosis of malaria with 96 and 100 per cent sensitivity and 100 per cent specificity when the parasite count was > 75 parasite/ μ l blood. Our data also provided a standardized protocol for the measurement of both *Pf*LDH and *Pf*HRP-2 antigens in the same patient sample using separate capture or detecting antibodies specific to these antigens.

In conclusion, both the antigen capture assays were comparable to each other for detection of *P. falciparum* infection at low levels of parasitaemia. The sensitivity limit (0.0015% parasitaemia) of PfHRP-2 and PfLDH assays was comparable to those achieved by microscopic examination of thick blood smears or other RDTs.

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