SUPPLEMENTAL MATERIAL

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Table S1. *Plasmodium* antigen-specific monoclonal antibodies used in screening for the 4-Plex array

 development

Biomarker	Clone	Host	Property	Ig class	Supplier	Applications
HRP2	MPFM-55A	Mouse	Monoclonal	IgM	Immunology consultants lab	ELISA, LF, WB
	MPFG-55A	Mouse	Monoclonal	IgG	Immunology consultants lab	ELISA, LF, WB
	PTL3	Mouse	Monoclonal	IgM	National Bioproducts Institute	ELISA, LF, WB
	C1-13	Mouse	Monoclonal	IgG	National Bioproducts Institute	ELISA, LF, WB
	0445	Mouse	Monoclonal	IgG	Vista Diagnostics	ELISA
	2G6	Mouse	Monoclonal	IgG	Vista Diagnostics	ELISA
	4D6	Mouse	Monoclonal	IgG	Precision Antibody	ELISA
	6C8	Mouse	Monoclonal	IgG	Precision Antibody	ELISA
	8D3	Mouse	Monoclonal	IgG	Precision Antibody	ELISA
	10C1	Mouse	Monoclonal	IgG	Precision Antibody	ELISA
	10F5	Mouse	Monoclonal	IgG	Precision Antibody	ELISA
	11E10	Mouse	Monoclonal	IgG	Precision Antibody	ELISA
	12D4	Mouse	Monoclonal	IgG	Precision Antibody	ELISA
	12F12	Mouse	Monoclonal	IgG	Precision Antibody	ELISA
Pan LDH	1201	Mouse	Monoclonal	IgG	Vista Diagnostics	ELISA
	19G7	Mouse	Monoclonal	IgG	Vista Diagnostics	ELISA
	1246	Mouse	Monoclonal	IgG	Vista Diagnostics	ELISA
	12G1	Mouse	Monoclonal	IgG	Vista Diagnostics	ELISA
	6G9	Mouse	Monoclonal	IgG	Vista Diagnostics	ELISA
P. vivax LDH	3H8	Mouse	Monoclonal	IgG	Vista Diagnostics	ELISA
	1102	Mouse	Monoclonal	IgG	Vista Diagnostics	ELISA
	13H4	Mouse	Monoclonal	IgG	Vista Diagnostics	ELISA

Abbreviation: ELISA, enzyme-linked immunosorbent assay; LF, lateral flow; WB, western blot

	Signal-to-noise ratio of 10 pg/mL to blank; HRP2													
Detection Capture	MPFM- 55A	MPFG- 55A	PTL3	C1-13	0445	2G6	4D6	6C8	8D3	10C1	10F5	11E10	12 D 4	12F12
MPFM-55A	1.4	15.3	1.3	6.3	3.1	5.1	2.5	16.9	4.2	4.9	25.6	40.4	26.5	5.6
MPFG-55A	4.3	5.5	1.9	5.8	2.8	8.5	1.4	8.3	1.5	4.6	16.5	22.4	8.7	4.7
PTL3	1.7	10.8	1.2	5.9	3.6	3.7	2.6	12.4	2.0	2.4	21.3	24.9	16.2	8.4
C1-13	3.9	7.2	1.5	4.9	2.3	4.5	2.0	5.1	1.1	2.9	13.2	14.0	5.7	5.6
445	5.0	5.7	1.4	5.4	3.1	14.5	3.1	9.5	1.3	4.0	20.2	31.5	13.3	5.9
2G6	2.4	12.7	1.0	2.9	3.0	8.2	1.5	18.7	4.0	2.1	41.7	55.3	10.4	8.2
4D6	2.5	2.5	1.0	3.2	2.0	8.0	1.0	6.0	2.1	1.0	9.9	20.6	1.2	1.9
6C8	1.0	1.3	0.9	1.1	1.3	3.4	0.9	1.5	1.2	1.0	1.5	4.2	1.0	0.9
8D3	3.9	6.0	1.3	5.0	2.4	14.2	2.2	7.2	1.1	2.5	18.3	25.3	3.7	4.4
10C1	3.5	3.9	0.9	4.0	2.0	8.5	1.5	6.4	3.9	2.4	14.8	33.3	4.4	4.5
10F5	2.5	2.3	0.9	3.3	2.0	7.3	2.4	7.5	2.6	2.8	15.2	25.1	8.1	5.4
11E10	1.2	1.5	1.0	1.1	1.5	2.7	1.0	2.3	2.4	1.0	1.2	2.5	1.0	1.0
12D4	3.3	3.4	1.4	4.4	2.3	8.9	1.2	7.6	2.4	1.8	19.2	26.0	1.7	3.0
12F12	3.0	3.1	1.1	4.0	1.9	6.7	1.5	9.5	2.5	2.1	17.2	32.8	3.2	5.2

Table S2. Rapid antibody pairing to identify the high performing antibody pair in detecting HRP2

For the rapid antibody pairing study, an array immobilized with 14 of anti-HRP2 monoclonal antibodies on the bottom of each well of 96-well plate was screened pairwise against 10, 100, and 1,000 pg/ml of recombinant HRP2 protein. The performance of each capture-detection pair was screened by measuring signal-to-noise ratio. Signal-to-noise ratio of 10 pg/ml to blank sample is shown. The heat map was created using the Quick Analysis tool from Excel. The red in color represents the stronger binding response whereas the blue for the poor binding response.

Table S3. Rapid antibody	pairing to identify the	high performing a	antibody pair in detection	ng Pan LDH and
P. vivax LDH				

	Signal-to-noise ratio of 1000 pg/ml to blank; P. falciparum LDH									
Detection Capture	1201	19G7	1246	12G1	6G9	3H8	1102	13H1		
1201	57.4	27.6	62.8	16.0	9.6	0.9	1.0	0.9		
19G7	133.3	2.4	39.9	4.0	16.8	1.0	1.1	1.0		
1246	38.1	12.6	47.5	1.5	8.4	0.9	1.0	1.0		
12G1	103.6	4.2	59.1	1.7	23.5	1.0	0.8	0.9		
6G9	16.7	6.3	19.9	3.7	3.1	1.0	0.9	1.0		
3H8	1.1	0.9	1.5	1.0	1.1	1.2	1.0	1.1		
1102	1.4	0.9	0.9	0.9	0.8	0.9	1.2	1.0		
13H1	1.0	1.2	1.3	0.9	0.8	1.1	0.8	1.2		
	Signal-to-noise ratio of 1000 pg/ml to blank; P. vivax LDH									
		s	ignal-to-noise	ratio of 1000	pg/ml to blank	; P. vivax LD	H			
Detection Capture	1201	S 19G7	ignal-to-noise 1246	ratio of 1000 12G1	pg/ml to blank 6G9	; <i>P. vivax</i> LD 3H8	H 1102	13H1		
Detection Capture 1201	1201 46.0	S 19G7 34.0	ignal-to-noise 1246 10.1	ratio of 1000 12G1 24.1	pg/ml to blank 6G9 1.4	; <i>P. vivax</i> LD 3H8 39.9	H 1102 56.2	13H1 9.0		
Detection Capture 1201 19G7	1201 46.0 113.1	S 19G7 34.0 4.0	ignal-to-noise 1246 10.1 8.8	ratio of 1000 12G1 24.1 12.6	pg/ml to blank 6G9 1.4 1.7	; <i>P. vivax</i> LD 3H8 39.9 37.7	H 1102 56.2 21.7	13H1 9.0 14.1		
Detection Capture 1201 19G7 1246	1201 46.0 113.1 42.1	\$ 19G7 34.0 4.0 14.7	ignal-to-noise 1246 10.1 8.8 6.0	ratio of 1000 12G1 24.1 12.6 2.2	pg/ml to blank 6G9 1.4 1.7 1.3	; <i>P. vivax</i> LD 3H8 39.9 37.7 17.1	H 1102 56.2 21.7 6.2	13H1 9.0 14.1 3.8		
Detection Capture 1201 19G7 1246 12G1	1201 46.0 113.1 42.1 169.3	S 19G7 34.0 4.0 14.7 5.5	ignal-to-noise 1246 10.1 8.8 6.0 13.5	ratio of 1000 12G1 24.1 12.6 2.2 2.0	pg/ml to blank 6G9 1.4 1.7 1.3 1.6	; <i>P. vivax</i> LD 3H8 39.9 37.7 17.1 30.5	H 1102 56.2 21.7 6.2 1.0	13H1 9.0 14.1 3.8 2.0		
Detection Capture 1201 19G7 1246 12G1 6G9	1201 46.0 113.1 42.1 169.3 9.8	S 19G7 34.0 4.0 14.7 5.5 4.7	ignal-to-noise 1246 10.1 8.8 6.0 13.5 2.2	ratio of 1000 12G1 24.1 12.6 2.2 2.0 4.2	pg/ml to blank 6G9 1.4 1.7 1.3 1.6 1.1	; <i>P. vivax</i> LD 3H8 39.9 37.7 17.1 30.5 6.3	H 1102 56.2 21.7 6.2 1.0 1.0	13H1 9.0 14.1 3.8 2.0 1.9		
Detection Capture 1201 19G7 1246 12G1 6G9 3H8	1201 46.0 113.1 42.1 169.3 9.8 41.6	S 19G7 34.0 4.0 14.7 5.5 4.7 4.7	ignal-to-noise 1246 10.1 8.8 6.0 13.5 2.2 3.3	ratio of 1000 12G1 24.1 12.6 2.2 2.0 4.2 2.1	pg/ml to blank 6G9 1.4 1.7 1.3 1.6 1.1 1.0	; P. vivax LD 3H8 39.9 37.7 17.1 30.5 6.3 2.1	H 1102 56.2 21.7 6.2 1.0 1.0 1.0	13H1 9.0 14.1 3.8 2.0 1.9 1.0		
Detection Capture 1201 19G7 1246 12G1 6G9 3H8 1102	1201 46.0 113.1 42.1 169.3 9.8 41.6 52.7	S 19G7 34.0 4.0 14.7 5.5 4.7 4.7 5.5	ignal-to-noise 1246 10.1 8.8 6.0 13.5 2.2 3.3 4.0	ratio of 1000 12G1 24.1 12.6 2.2 2.0 4.2 2.1 1.2	pg/ml to blank 6G9 1.4 1.7 1.3 1.6 1.1 1.0 0.9	; P. vivax LD 3H8 39.9 37.7 17.1 30.5 6.3 2.1 1.1	H 1102 56.2 21.7 6.2 1.0 1.0 1.0 4.2	13H1 9.0 14.1 3.8 2.0 1.9 1.0 1.3		

For the rapid antibody pairing study, an array immobilized with 8 of anti-pan LDH and anti-*P. vivax* LDH monoclonal antibodies on the bottom of each well of 96-well plate was screened pairwise against 100, 1,000, and 10,000 pg/ml of either *P. falciparum* LDH or *P. vivax* LDH proteins. The performance of each capture-detection pair was screened by measuring signal-to-noise ratio. Signal-to-noise ratio of 1,000 pg/ml to blank sample is shown. The heat map was created using the Quick Analysis tool from Excel. The red in color represents the stronger binding response whereas the blue for the poor binding response.

Biomarker	Capture antibody	Detection antibody	Calibrato	Assay
HRP2	0445 (Vista)	11E10 (Precision)	HRP2 (Microcoat)	Quantitative
Pan LDH	12G1 (Vista)	1201 (Vista)	P. vivax LDH (CTK)	Qualitative ^b /Quantitative
P. vivax LDH	1102 (Vista)	1201 (Vista)	P. vivax LDH (CTK)	Quantitative
Human CRP	C6 (HyTest)	n.a ª	CRP (HyTest)	Quantitative

Table S4. Final monoclonal antibodies and calibrator proteins with their sources used on the 4-Plex array

^a Quantification of human CRP is based on competitive ELISA using biotinylated CRP. Supplier is shown in parenthesis.

^b Pan LDH assay may be used for quantification of pLDH in a specimen infected with *P. falciparum* species.

HRP2 (pg/mL)	Pixel inte Mean, CV	ensity V%	Back-ca Mean, C	lculated V%	Accuracy % ^a	Pan LDH (pg/mL)	Pixel intensity Mean, CV%	Back-calculated Mean, CV%	Accuracy %ª
590	47833.3	1.7	588.0	28.5	99.7	10,514	33431.5 5.7	10596.0 11.2	100.8
196.7	41176.1	2.5	190.8	8.0	97.0	3,504.7	15460.0 9.3	3480.3 10.5	99.3
65.6	22433.3	6.9	66.8	8.2	101.8	1,168.2	5571.2 8.2	1155.5 9.0	98.9
21.9	7747.8	8.5	21.6	8.5	98.8	389.4	2023.5 7.9	390.8 9.1	100.4
7.3	2692.3	9.7	7.5	9.9	102.3	129.8	798.9 8.8	128.7 11.3	99.1
2.4	907.7	16.9	2.3	20.2	93.7	43.3	388.8 7.8	42.2 16.0	97.5
0.8	450.2	14.9	0.9	23.8	109.0	14.4	259.5 6.9	14.4 26.8	100.1
0.3	252.7	25.0	0.3	72.4	101.4	4.8	213.2 13.9	7.3 65.8	151.7
P. vivax LDH	Pixel int	ensity	Back-ca	alculated		CRP	Pixel intensity	Back-calculated	
					A a a 11 m a a 11 0 / 2	Cit	2		A = 0/3
(pg/mL)	Mean, C	V%	Mean, C	CV%	Accuracy % ^a	(ng/mL)	Mean, CV%	Mean, CV%	Accuracy % ^a
(pg/mL) 	Mean, C 10365.1	V%	Mean, 0 479.6	2V% 9.9	Accuracy % ^a 96.5	(ng/mL) 13,500	Mean, CV%	Mean, CV% 13713.3 11.2	Accuracy % ^a 101.6
(pg/mL) 497.0 165.7	Mean, C 10365.1 3990.5	8.2 9.1	Mean, C 479.6 165.5	2V% 9.9 10.0	Accuracy % ^a 96.5 99.9	(ng/mL) 13,500 4,500	Mean, CV% 496.7 9.4 1278.0 12.3	Mean, CV% 13713.3 11.2 4479.7 16.8	Accuracy % ^a 101.6 99.5
(pg/mL) 497.0 165.7 55.2	Mean, C 10365.1 3990.5 1457.0	8.2 9.1 8.1	Mean, C 479.6 165.5 54.7	9.9 10.0 9.9	Accuracy % ^a 96.5 99.9 99.0	(ng/mL) 13,500 4,500 1,500	Mean, CV% 496.7 9.4 1278.0 12.3 3109.5 11.0	Mean, CV% 13713.3 11.2 4479.7 16.8 1543.4 13.8	Accuracy % ^a 101.6 99.5 102.9
(pg/mL) 497.0 165.7 55.2 18.4	Mean, C 10365.1 3990.5 1457.0 637.1	8.2 9.1 8.1 6.2	Mean, C 479.6 165.5 54.7 18.7	9.9 10.0 9.9 9.7	Accuracy % ^a 96.5 99.9 99.0 101.4	(ng/mL) 13,500 4,500 1,500 500	Mean, CV% 496.7 9.4 1278.0 12.3 3109.5 11.0 7732.9 10.2	Mean, CV% 13713.3 11.2 4479.7 16.8 1543.4 13.8 505.4 13.1	Accuracy % ^a 101.6 99.5 102.9 101.1
(pg/mL) 497.0 165.7 55.2 18.4 6.1	Mean, C 10365.1 3990.5 1457.0 637.1 347.8	8.2 9.1 8.1 6.2 8.0	Mean, C 479.6 165.5 54.7 18.7 6.1	9.9 10.0 9.9 9.7 20.6	Accuracy % ^a 96.5 99.9 99.0 101.4 99.1	(ng/mL) 13,500 4,500 1,500 500 166.7	Mean, CV% 496.7 9.4 1278.0 12.3 3109.5 11.0 7732.9 10.2 16965.8 10.2	Mean, CV% 13713.3 11.2 4479.7 16.8 1543.4 13.8 505.4 13.1 159.8 18.8	Accuracy % ^a 101.6 99.5 102.9 101.1 95.9
(pg/mL) 497.0 165.7 55.2 18.4 6.1 2.1	Mean, C 10365.1 3990.5 1457.0 637.1 347.8 251.0	8.2 9.1 8.1 6.2 8.0 10.2	Mean, C 479.6 165.5 54.7 18.7 6.1 2.4	9.9 10.0 9.9 9.7 20.6 29.6	Accuracy % ^a 96.5 99.9 99.0 101.4 99.1 115.1	(ng/mL) 13,500 4,500 1,500 500 166.7 55.6	Mean, CV% 496.7 9.4 1278.0 12.3 3109.5 11.0 7732.9 10.2 16965.8 10.2 24216.3 6.2	Mean, CV% 13713.3 11.2 4479.7 16.8 1543.4 13.8 505.4 13.1 159.8 18.8 61.5 28.7	Accuracy % ^a 101.6 99.5 102.9 101.1 95.9 110.6
(pg/mL) 497.0 165.7 55.2 18.4 6.1 2.1 0.7	Mean, C 10365.1 3990.5 1457.0 637.1 347.8 251.0 223.0	8.2 9.1 8.1 6.2 8.0 10.2 14.2 14.2	Mean, C 479.6 165.5 54.7 18.7 6.1 2.4 1.3	9.9 10.0 9.9 9.7 20.6 29.6 81.3	Accuracy % ^a 96.5 99.9 99.0 101.4 99.1 115.1 193.5	(ng/mL) 13,500 4,500 1,500 500 166.7 55.6 18.5	Mean, CV% 496.7 9.4 1278.0 12.3 3109.5 11.0 7732.9 10.2 16965.8 10.2 24216.3 6.2 27828.5 9.0	Mean, CV% 13713.3 11.2 4479.7 16.8 1543.4 13.8 505.4 13.1 159.8 18.8 61.5 28.7 37.3 73.6	Accuracy % ^a 101.6 99.5 102.9 101.1 95.9 110.6 201.2

Table S5. Characterization of calibrator curves generated during assay validation

^a Accuracy was calculated using the formula; observed concentration/expected concentration) x 100. Data were analyzed from twelve calibration curves obtained on different days.

Expected conc. (pg/mL)	Mean adjusted				
	conc. (pg/mL)	Accuracy %	Expected conc. (pg/mL)	Mean adjusted conc. (pg/mL)	Accuracy %
45.0	38.2	84.9	55.0	53.2	96.8
22.5	22.1	98.2	27.5	27.8	101.1
11.3	11.2	99.2	13.8	<lloq< td=""><td></td></lloq<>	
5.6	5.8	102.6	6.9	<lloq< td=""><td></td></lloq<>	
2.8	3.0	106.5	3.4	<lloq< td=""><td></td></lloq<>	
1.4	1.6	112.0	1.7	<lloq< td=""><td></td></lloq<>	
325.0	332.3	102.2	300.0	267.0	89.0
162.5	167.2	102.9	150.0	140.1	93.4
81.3	80.1	98.5	75.0	72.1	96.2
40.6	39.5	97.1	37.5	37.9	101.1
20.3	19.5	95.9	18.8	18.9	100.7
10.2	10.1	99.5	9.4	<lloq< td=""><td></td></lloq<>	
275.0	318.7	115.9	200.0	174.4	87.2
137.5	135.5	98.5	100.0	95.9	95.9
68.8	65.9	95.9	50.0	51.8	103.6
34.4	32.1	93.3	25.0	25.0	99.8
17.2	16.3	94.8	12.5	<lloq< td=""><td></td></lloq<>	
8.6	8.6	100.2	6.3	<lloq< td=""><td></td></lloq<>	
ND	<lloq< td=""><td></td><td>875.0</td><td>831.5</td><td>95.0</td></lloq<>		875.0	831.5	95.0
ND	<lloq< td=""><td></td><td>437.5</td><td>446.0</td><td>101.9</td></lloq<>		437.5	446.0	101.9
ND	<lloq< td=""><td></td><td>218.8</td><td>215.5</td><td>98.5</td></lloq<>		218.8	215.5	98.5
ND	<lloq< td=""><td></td><td>109.4</td><td>110.0</td><td>100.6</td></lloq<>		109.4	110.0	100.6
ND	<lloq< td=""><td></td><td>54.7</td><td>55.6</td><td>101.6</td></lloq<>		54.7	55.6	101.6
ND	<lloq< td=""><td></td><td>27.3</td><td>29.5</td><td>107.8</td></lloq<>		27.3	29.5	107.8
ND	<lloq< td=""><td></td><td>1040.7</td><td>1050.0</td><td>99.1</td></lloq<>		1040.7	1050.0	99.1
ND	<lloq< td=""><td></td><td>532.0</td><td>525.0</td><td>101.3</td></lloq<>		532.0	525.0	101.3
ND	<lloq< td=""><td></td><td>253.3</td><td>262.5</td><td>96.5</td></lloq<>		253.3	262.5	96.5
ND	<lloq< td=""><td></td><td>129.1</td><td>131.3</td><td>98.4</td></lloq<>		129.1	131.3	98.4
ND	<lloq< td=""><td></td><td>62.2</td><td>65.6</td><td>94.8</td></lloq<>		62.2	65.6	94.8
ND	<lloq< td=""><td></td><td>38.0</td><td>32.8</td><td>115.9</td></lloq<>		38.0	32.8	115.9
	ND ND ND ND ND	ND <lloq ND <lloq ND <lloq ND <lloq ND <lloq ND <lloq< td=""><td>ND <lloq ND <lloq ND <lloq ND <lloq ND <lloq ND <lloq< td=""><td>ND <lloq< th=""> 1040.7 ND <lloq< td=""> 532.0 ND <lloq< td=""> 253.3 ND <lloq< td=""> 129.1 ND <lloq< td=""> 62.2 ND <lloq< td=""> 38.0</lloq<></lloq<></lloq<></lloq<></lloq<></lloq<></td><td>ND <lloq< th=""> 1040.7 1050.0 ND <lloq< td=""> 532.0 525.0 ND <lloq< td=""> 253.3 262.5 ND <lloq< td=""> 129.1 131.3 ND <lloq< td=""> 62.2 65.6 ND <lloq< td=""> 38.0 32.8</lloq<></lloq<></lloq<></lloq<></lloq<></lloq<></td></lloq<></lloq </lloq </lloq </lloq </lloq </td></lloq<></lloq </lloq </lloq </lloq </lloq 	ND <lloq ND <lloq ND <lloq ND <lloq ND <lloq ND <lloq< td=""><td>ND <lloq< th=""> 1040.7 ND <lloq< td=""> 532.0 ND <lloq< td=""> 253.3 ND <lloq< td=""> 129.1 ND <lloq< td=""> 62.2 ND <lloq< td=""> 38.0</lloq<></lloq<></lloq<></lloq<></lloq<></lloq<></td><td>ND <lloq< th=""> 1040.7 1050.0 ND <lloq< td=""> 532.0 525.0 ND <lloq< td=""> 253.3 262.5 ND <lloq< td=""> 129.1 131.3 ND <lloq< td=""> 62.2 65.6 ND <lloq< td=""> 38.0 32.8</lloq<></lloq<></lloq<></lloq<></lloq<></lloq<></td></lloq<></lloq </lloq </lloq </lloq </lloq 	ND <lloq< th=""> 1040.7 ND <lloq< td=""> 532.0 ND <lloq< td=""> 253.3 ND <lloq< td=""> 129.1 ND <lloq< td=""> 62.2 ND <lloq< td=""> 38.0</lloq<></lloq<></lloq<></lloq<></lloq<></lloq<>	ND <lloq< th=""> 1040.7 1050.0 ND <lloq< td=""> 532.0 525.0 ND <lloq< td=""> 253.3 262.5 ND <lloq< td=""> 129.1 131.3 ND <lloq< td=""> 62.2 65.6 ND <lloq< td=""> 38.0 32.8</lloq<></lloq<></lloq<></lloq<></lloq<></lloq<>

Table S6. Dilutional linearity with the 4-Plex array

			P. vivax LDH			CRP	
		Expected conc. (pg/mL)	Mean adjusted conc. (pg/mL)	Accuracy %	Expected conc. (pg/mL)	Mean adjusted conc. (pg/mL)	Accuracy %
Sample 1	1:4	ND	<lloq< td=""><td></td><td>ND</td><td><lloq< td=""><td></td></lloq<></td></lloq<>		ND	<lloq< td=""><td></td></lloq<>	
	1:8	ND	<lloq< td=""><td></td><td>ND</td><td><lloq< td=""><td></td></lloq<></td></lloq<>		ND	<lloq< td=""><td></td></lloq<>	
	1:16	ND	<lloq< td=""><td></td><td>ND</td><td><lloq< td=""><td></td></lloq<></td></lloq<>		ND	<lloq< td=""><td></td></lloq<>	
	1:32	ND	<lloq< td=""><td></td><td>ND</td><td><lloq< td=""><td></td></lloq<></td></lloq<>		ND	<lloq< td=""><td></td></lloq<>	
	1:64	ND	<lloq< td=""><td></td><td>ND</td><td><lloq< td=""><td></td></lloq<></td></lloq<>		ND	<lloq< td=""><td></td></lloq<>	
	1:128	ND	<lloq< td=""><td></td><td>ND</td><td><lloq< td=""><td></td></lloq<></td></lloq<>		ND	<lloq< td=""><td></td></lloq<>	
ample 2	1:4	ND	<lloq< td=""><td></td><td>70.0</td><td>69.5</td><td>99.2</td></lloq<>		70.0	69.5	99.2
	1:8	ND	<lloq< td=""><td></td><td>35.0</td><td><lloq< td=""><td></td></lloq<></td></lloq<>		35.0	<lloq< td=""><td></td></lloq<>	
	1:16	ND	<lloq< td=""><td></td><td>17.5</td><td><lloq< td=""><td></td></lloq<></td></lloq<>		17.5	<lloq< td=""><td></td></lloq<>	
	1:32	ND	<lloq< td=""><td></td><td>8.8</td><td><lloq< td=""><td></td></lloq<></td></lloq<>		8.8	<lloq< td=""><td></td></lloq<>	
	1:64	ND	<lloq< td=""><td></td><td>4.4</td><td><lloq< td=""><td></td></lloq<></td></lloq<>		4.4	<lloq< td=""><td></td></lloq<>	
	1:128	ND	<lloq< td=""><td></td><td>2.2</td><td><lloq< td=""><td></td></lloq<></td></lloq<>		2.2	<lloq< td=""><td></td></lloq<>	
ample 3	1:4	ND	<lloq< td=""><td></td><td>450.0</td><td>495.3</td><td>110.1</td></lloq<>		450.0	495.3	110.1
-	1:8	ND	<lloq< td=""><td></td><td>225.0</td><td>200.7</td><td>89.2</td></lloq<>		225.0	200.7	89.2
	1:16	ND	<lloq< td=""><td></td><td>112.5</td><td><lloq< td=""><td></td></lloq<></td></lloq<>		112.5	<lloq< td=""><td></td></lloq<>	
	1:32	ND	<lloq< td=""><td></td><td>56.3</td><td><lloq< td=""><td></td></lloq<></td></lloq<>		56.3	<lloq< td=""><td></td></lloq<>	
	1:64	ND	<lloq< td=""><td></td><td>28.1</td><td><lloq< td=""><td></td></lloq<></td></lloq<>		28.1	<lloq< td=""><td></td></lloq<>	
	1:128	ND	<lloq< td=""><td></td><td>14.1</td><td><lloq< td=""><td></td></lloq<></td></lloq<>		14.1	<lloq< td=""><td></td></lloq<>	
ample 4	1:4	223.3	230.6	103.3	100.0	103.6	103.6
-	1:8	111.6	120.1	107.6	50.0	<lloq< td=""><td></td></lloq<>	
	1:16	55.8	54.4	97.5	25.0	<lloq< td=""><td></td></lloq<>	
	1:32	27.9	28.1	100.6	12.5	<lloq< td=""><td></td></lloq<>	
	1:64	13.9	13.2	94.8	6.3	<lloq< td=""><td></td></lloq<>	
	1:128	7.0	7.1	102.5	3.1	<lloq< td=""><td></td></lloq<>	
ample 5	1:4	266.0	296.4	111.4	ND	<lloq< td=""><td></td></lloq<>	
-	1:8	133.0	145.2	109.2	ND	<lloq< td=""><td></td></lloq<>	
	1:16	66.5	65.2	98.0	ND	<lloq< td=""><td></td></lloq<>	
	1:32	33.3	31.4	94.4	ND	<lloq< td=""><td></td></lloq<>	
	1:64	16.6	14.8	89.0	ND	<lloq< td=""><td></td></lloq<>	
	1:128	8.3	8.1	97.9	ND	<lloq< td=""><td></td></lloq<>	

^a Accuracy was calculated using the analyte concentration determined at dilution 1:4 as a reference value.

ND: not detected

Biomarker	No. of specimen ^a	Mean	Min	25 percentile	Median	75 percentile	Max
HRP2	39	7.295	0.001	0.008	0.049	0.386	108
Pan LDH	30	0.066	0.013	0.030	0.035	0.074	0.269
P. vivax LDH	41	0.006	0.001	0.002	0.003	0.008	0.038

Table S7. Descriptive statistics for amount of circulating antigen per parasite

a

^a Analysis was performed with specimens with parasitemia in the range of 1 to 100 parasites/ μ L and analyte concentration greater than the assay cutoff.



Fig S1. Concentration of HRP2 and Pan LDH, and parasitemia in blood specimens from all subjects obtained during a course of IBSM study. (A) concentration of HRP2 and Pan LDH, and (B) Parasitemia. The dotted lines indicate cut-off values.



Fig S2. Distribution of CRP levels in malaria non-infected and infected groups. Box plots of CRP levels in infection positive and negative groups determined by A) qPCR and B) 4-Plex array. Uninfected group determined by 4-Plex array was defined by the absence of any of malaria antigens. CRP levels were analyzed only from specimens which had quantifiable amount of HRP2, and *P. vivax* LDH greater than the assay cutoff. The results show the median (horizontal line), the 25th and 75the percentiles (boxes), the 10th and 90th percentiles (error bars) and the outliers (open circles). The two-tailed Mann–Whitney U-test was used to compare the distributions of two unmatched groups.