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THE SEARCH FOR

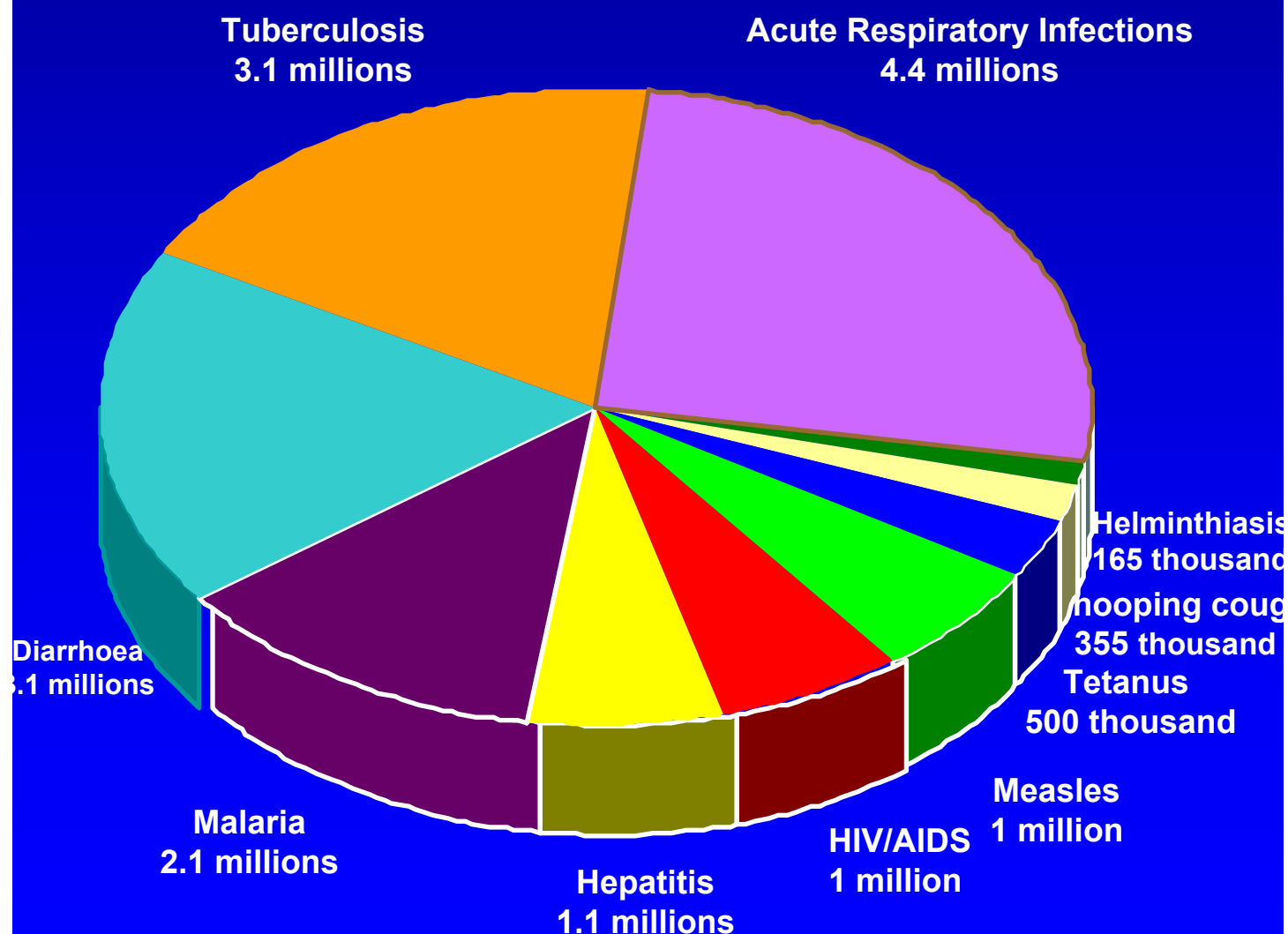
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Or

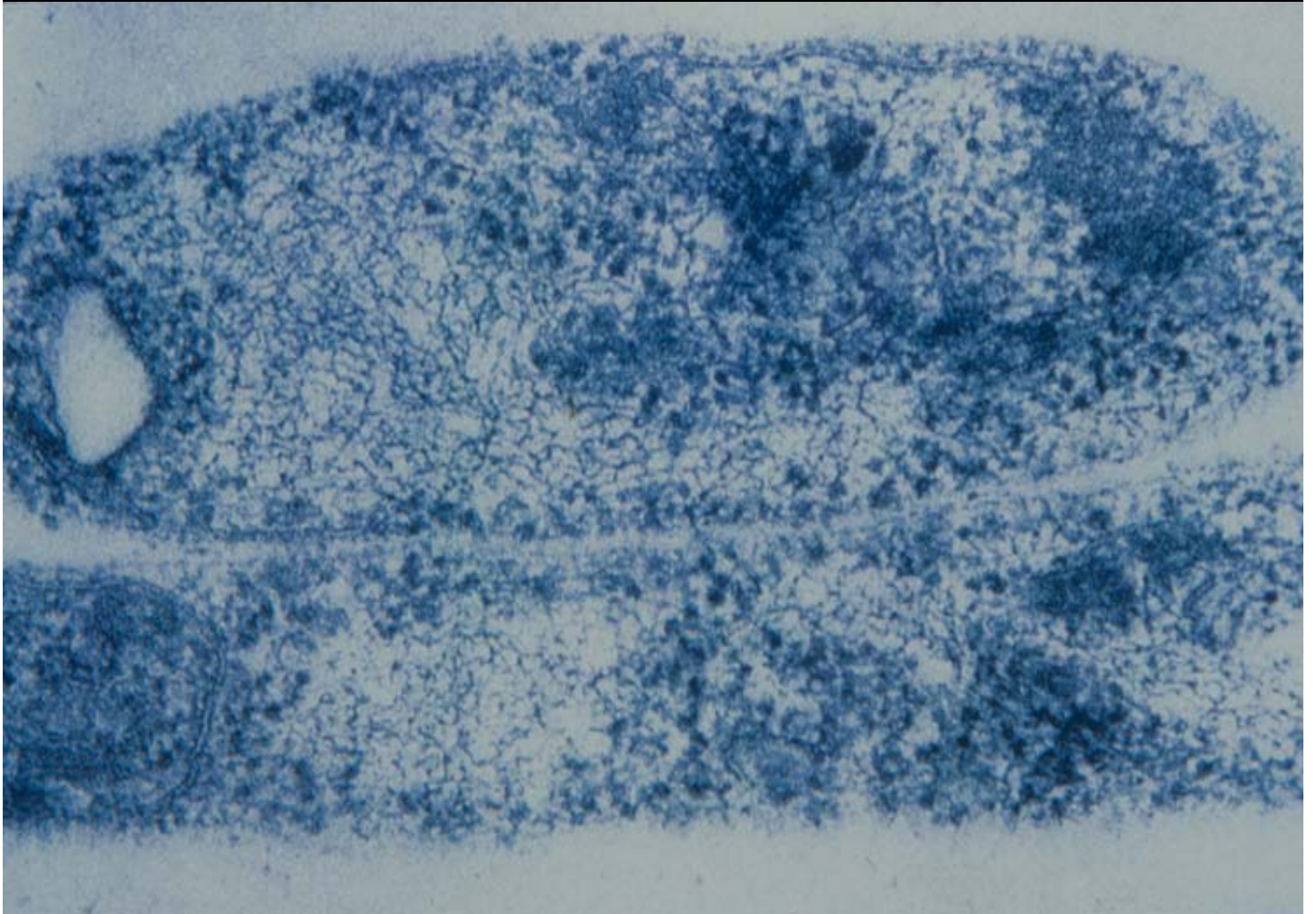
**A MATHEMATICAL
METHOD FOR
DEVELOPING
VACCINES**

TEN LEADING CAUSES OF MORTALITY IN THE WORLD INDUCED BY TRANSMITTABLE DISEASES

WHO-1995



BIOLOGICAL VACCINES



ACTUAL VACCINES

**SMALL POX
POLIO
MEASLES
PAROTIDITIS
GERMAN MEASLES
CHICKEN POX**

**HEPATITIS A & B
YELLOW FEVER
RABIES
BCG
DPT**

VACCINES IN PROGRESS

**MALARIA
AIDS
TYPHUS
CHOLERA
MENINGITIS
DENGUE
HERPES
S.T.D.
LEISHMANIA**

**SCHISTO
CARIES
LYME
STREPTOCOCCUS
RICKETS
INFLUENZA
ROTAVIRUS
RSV
PAPILOMA, ETC**

PURPOSE :

**To find a
rational approach
for the
development of
chemically
synthesized
vaccines.**

10 April 1997

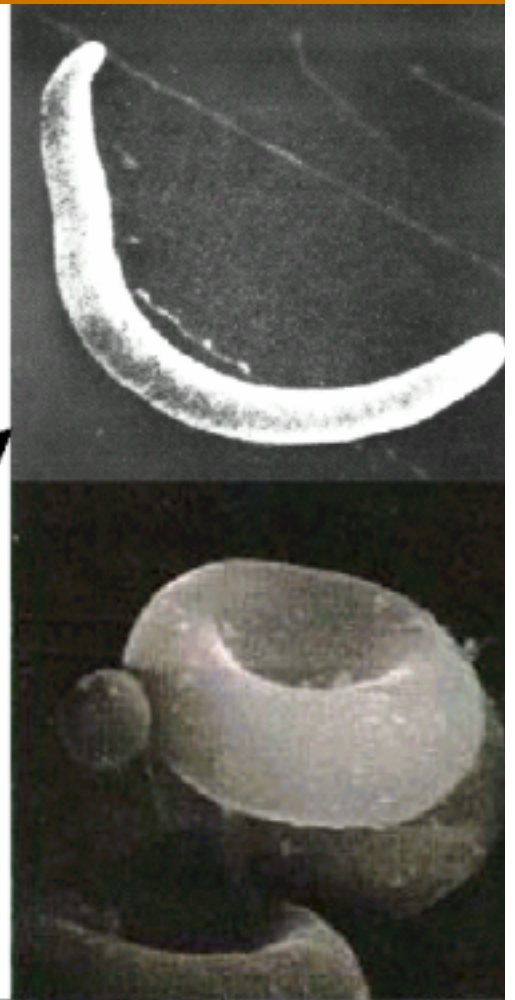
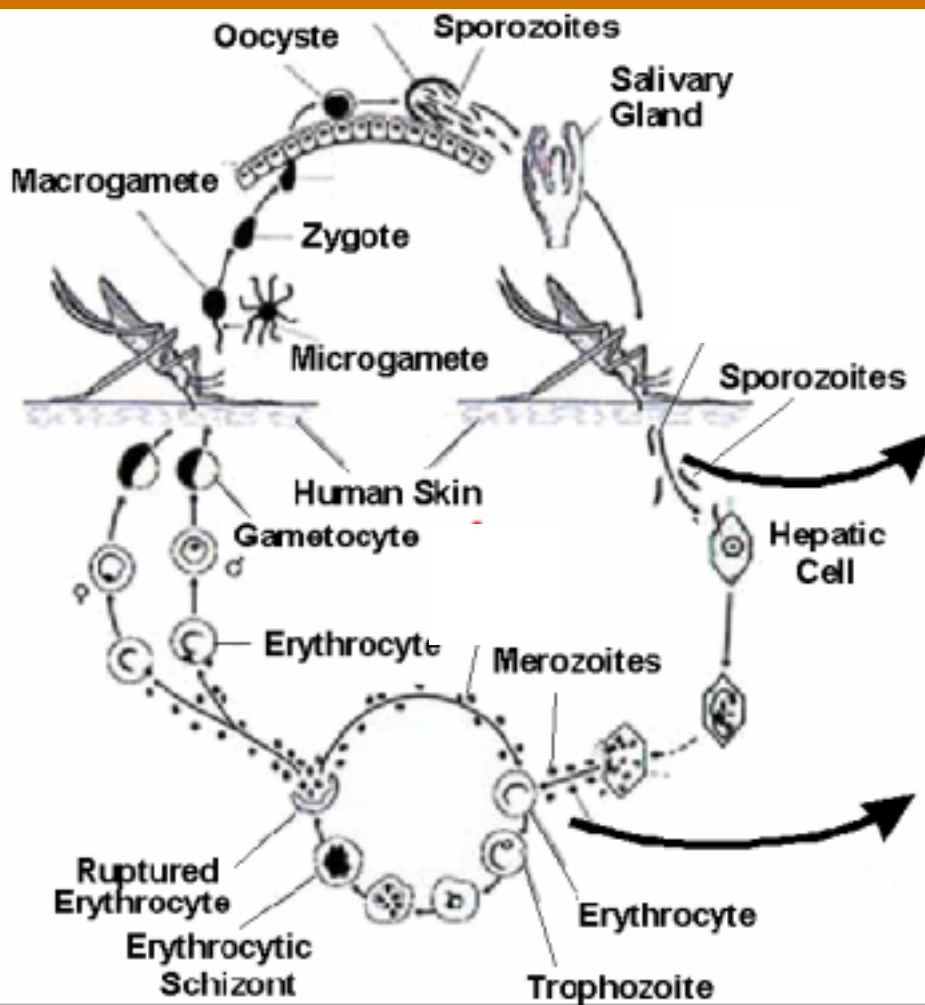
International weekly journal of science

nature

ISSN 0028-280X



Plasmodium falciparum PARASITE'S LIFE CYCLE



Why malaria?

- **300 million people are infected and 3 million die annually**
- **Acute disease**
- **Easy diagnosis**
- **Curable with treatment**
- **Experimental model: the Aotus monkey**
- **Heavy economic burden for the developing world**





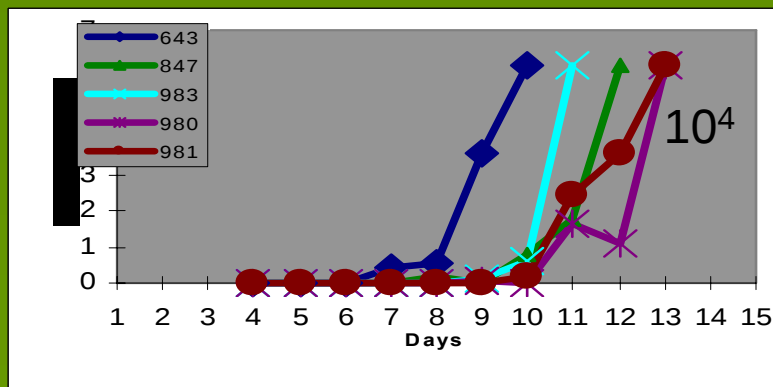
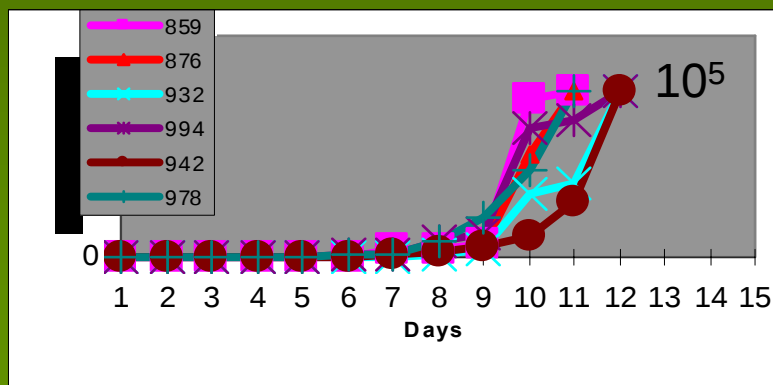
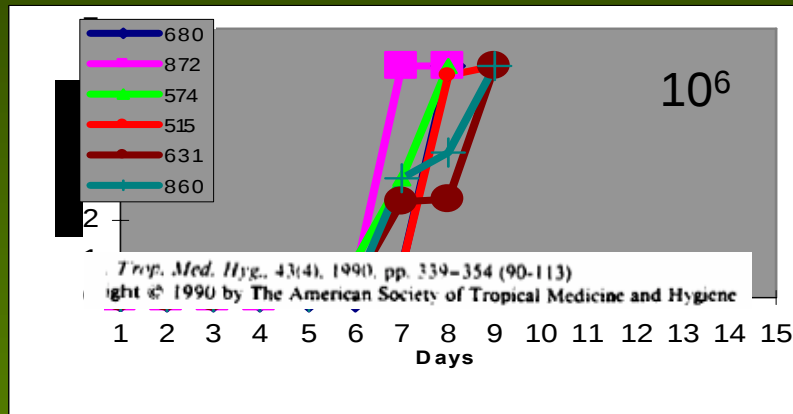
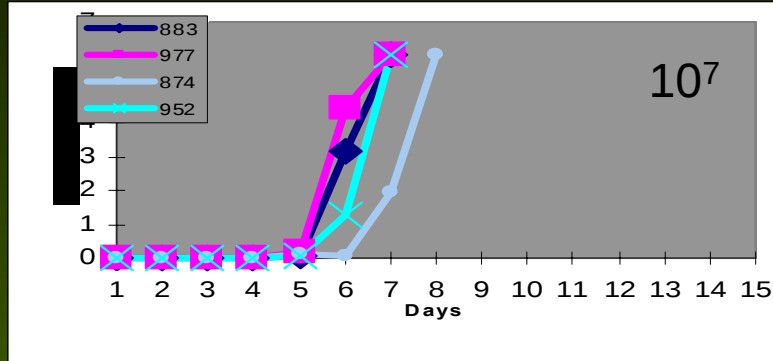






TITRATION OF THE FVO P. *falciparum* STRAIN IN AOTUS MONKEYS

Rodriguez R. et al. 1990. *Am. J. Trop. Med. Hyg* 43:339





ORIGINAL PAPER

Olga L. Díaz · Claudia A. Daubenberger
Raul Rodríguez · Martín Naegeli · Alberto Moreno

Immunoglobulin kappa light-chain *V*, *J*, and *C* gene sequences of the owl monkey *Aotus nancymae*

Identification and sequencing of genes from constant and variable regions (VH3 and VH4 families) of the heavy-chain of Immunoglobulin in Owl monkeys (*Aotus nancymae*)

EC HERNÁNDEZ, CF SUÁREZ, CA PARRA, OL DÍAZ AND ME PATARROYO.
Fundación Instituto de Inmunología de Colombia (FIDIC). Bogotá- Colombia. Calle 50 No. 26-00.

Immunogenetics (2002) 54:645–653
DOI 10.1007/s00251-002-0512-2

ORIGINAL PAPER

Edith C. Hernández · Carlos F. Suárez
Jairo A. Méndez · Sandra J. Echeverry
Luis A. Murillo · Manuel E. Patarroyo

Identification, cloning, and sequencing of different cytokine genes in four species of owl monkey

American Journal of Primatology 57:1–11 (2002)

RESEARCH ARTICLE

Partial Characterization of the CD45 Phosphatase DNA in the Owl Monkey (*Aotus vociferans*)

GLADIS E. MONTOYA¹, JEAN-PAUL VERNOT^{1,2*}, and MANUEL E. PATARROYO^{1,2}
¹Fundación Instituto de Inmunología de Colombia, FIDIC, Bogotá, D.C., Colombia
²Facultad de Medicina, Universidad Nacional de Colombia, Bogotá, D.C., Colombia

J Med Primatol 2003; 32: 31–38

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JOURNAL OF MEDICAL PRIMATOLOGY
ISSN 0047-2565

Proliferative response of peripheral blood lymphocytes to mitogens in the owl monkey *Aotus nancymae*

Pinzón-Charry A, Vernot JP, Rodríguez R, Patarroyo ME. Proliferative response of peripheral blood lymphocytes to mitogens in the owl monkey *Aotus nancymae*. *J med Primatol* 2003; 32:31–38. © Blackwell Munksgaard, 2003

Alberto Pinzón-Charry¹, Jean Paul Vernot^{1,2}, Raul Rodríguez¹, Manuel Elkin Patarroyo^{1,2}

C. F. Suárez*
P. P. Cárdenas*
E.J. Llanos-Ballesteras
P. Martínez
M. Obregón
M.E. Patarroyo
M.A. Patarroyo

**α_1 and α_2 domains of *Aotus* MHC Class I
and Catarrhini MHC Class Ia share similar
characteristics**

Tissue Antigens 2003: **61**: 362–373

Accepted in IMMUNOGENETICS
Unusual Variability in *Aotus* monkey MHC-E sequences

Paula P. Cárdenas¹, Carlos F. Suarez¹, Consuelo P.
Martínez, Eugenio J. Llanos, Mateo Obregón, Manuel E.
Patarroyo, Manuel A. Patarroyo *

J. E. Guerrero
D. P. Pacheco
C. F. Suárez
P. Martínez
F. Aristizabal
C. A. Moncada
M. E. Patarroyo
M. A. Patarroyo

**Characterizing T-cell receptor γ -variable
gene in *Aotus nancymae* owl monkey
peripheral blood**

Tissue Antigens 2003 **61**: 1–11

**Functional and Structural Similarity of $V\gamma 9V\delta 2$ T Cells in
Humans and *Aotus* Monkeys, a Primate Infection Model for
Plasmodium falciparum Malaria^{1,2}.**

The Journal of Immunology, 2001, 167: 6421–6430.

Claudia A. Daubenberger,^{3*} Maxence Salomon,* William Vecino,[†] Beatrice Hübner,*
Heike Troll,[‡] Raul Rodriques,[†] Manuel E. Patarroyo,[†] and Gerd Pluschke*

ORIGINAL PAPER

J. Javier Nino-Vasquez · Denise Vogel
Raul Rodriguez · Alberto Moreno
Manuel Elkin Patarroyo · Gerd Pluschke
Claudia A. Daubenberger

**Sequence and diversity of *DRB* genes of *Aotus nancymae*,
a primate model for human malaria parasites**

Immunogenetics (2000) 51: 528–537

Digital Object Identifier (DOI) 10.1007/s002510000189

ORIGINAL PAPER

Diana Diaz · Martin Naegeli · Raul Rodriguez
J. Javier Nino-Vasquez · Alberto Moreno
Manuel Elkin Patarroyo · Gerd Pluschke
Claudia A. Daubenberger

**Sequence and diversity of MHC *DQA* and *DQB* genes of the owl
monkey *Aotus nancymae***

Immunogenetics (2002) 54:251–259
DOI 10.1007/s00251-002-0466-4

ORIGINAL PAPER

Diana Diaz · Claudia A. Daubenberger · Tatjana Zalac
Raul Rodriguez · Manuel Elkin Patarroyo

**Sequence and expression of *MHC-DPB1* molecules
of the New World monkey *Aotus nancymae*, a primate model
for *Plasmodium falciparum***

ORIGINAL PAPER

Nicolas Favre · Claudia Daubenberger · Jutta Marfurt
Alberto Moreno · Manuel Patarroyo · Gerd Pluschke

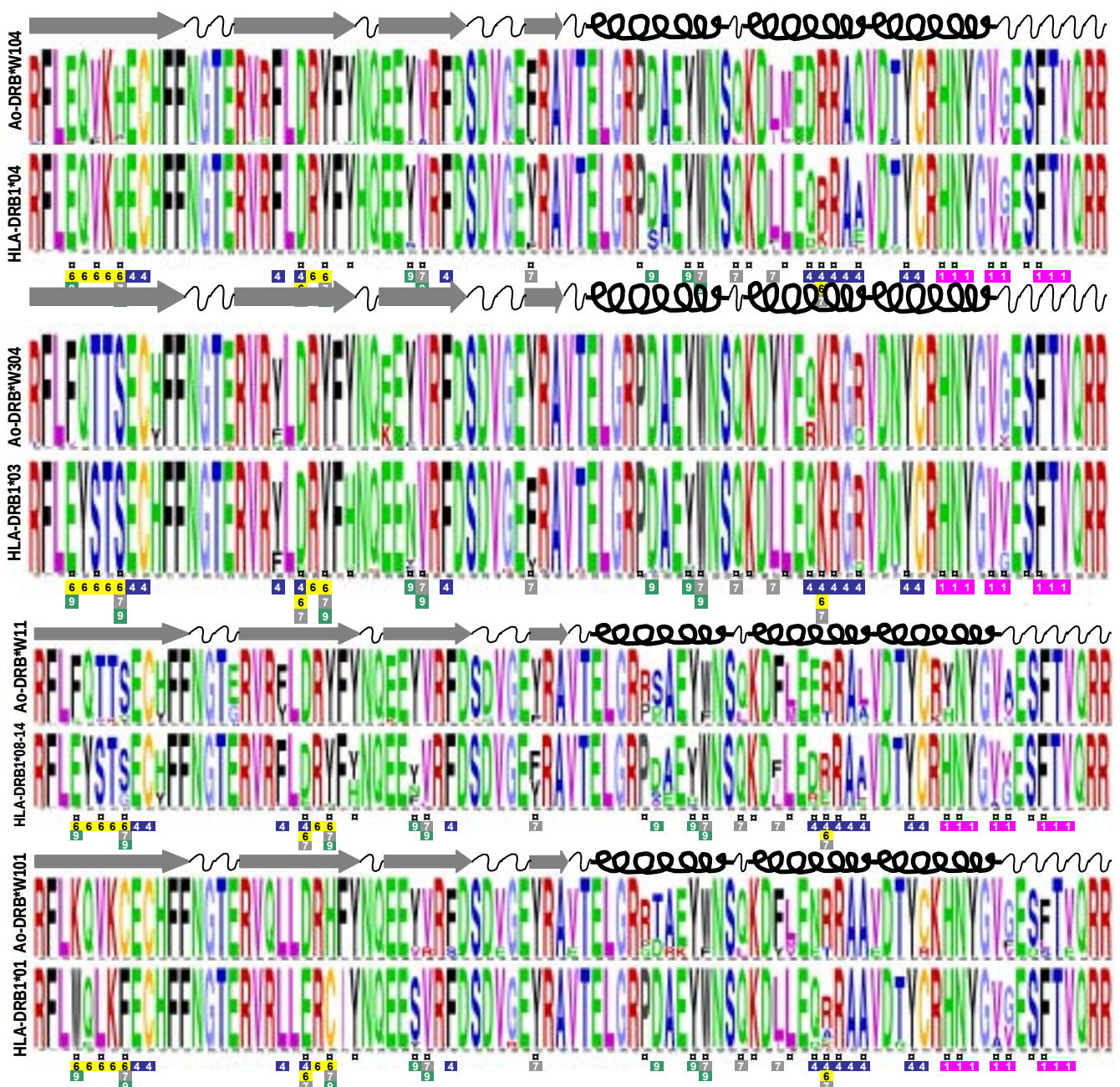
**Sequence and diversity of *T-cell receptor alpha* *V*, *J*, and *C* genes of
the owl monkey *Aotus nancymae***

ORIGINAL PAPER

William Vecino · Claudia Daubenberger
Raul Rodriguez · Alberto Moreno
Manuel Patarroyo · Gerd Pluschke

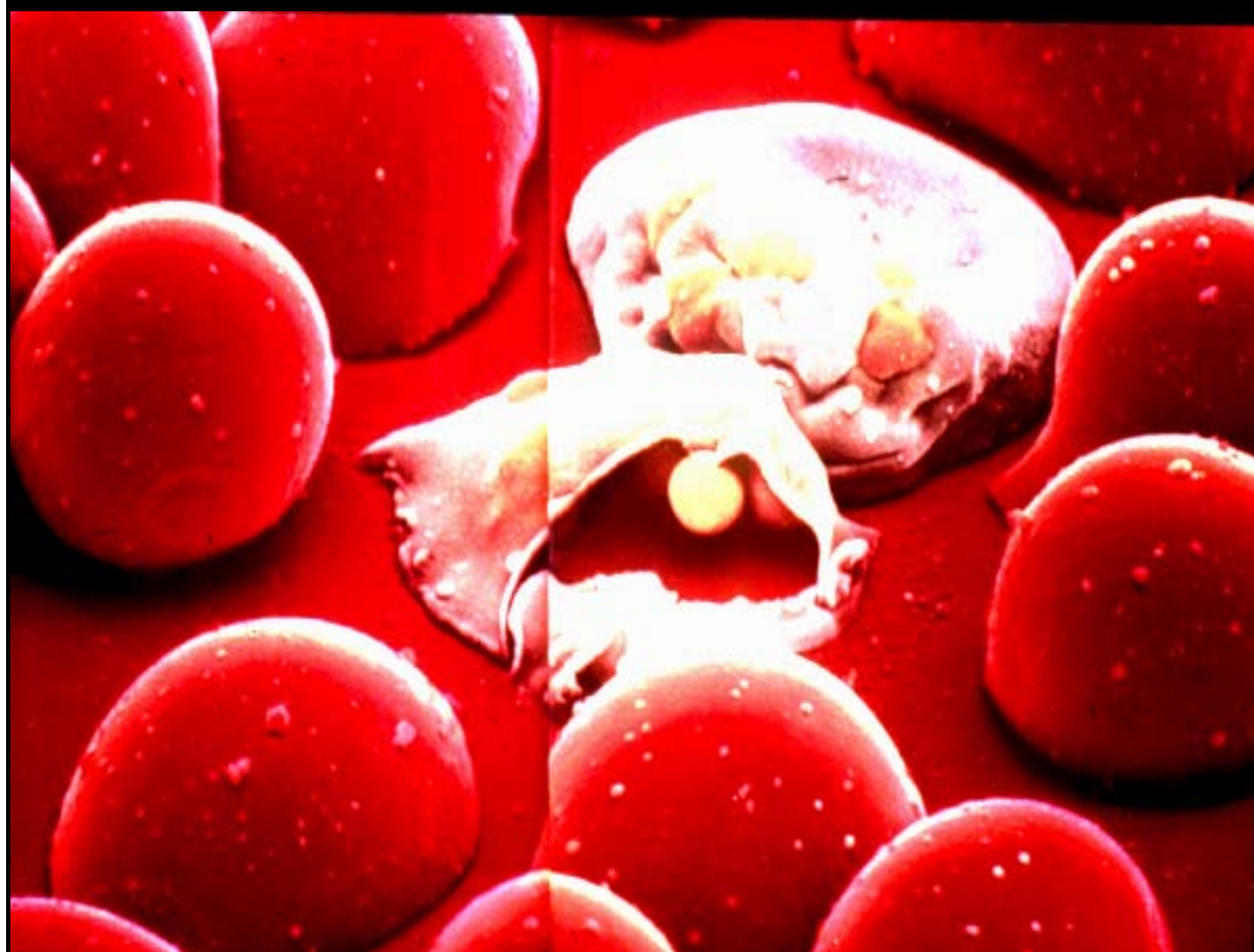
**Sequence and diversity of *T-cell receptor β -chain* *V* and *J* genes of the
owl monkey *Aotus nancymae***

Logos obtenidos a partir de las secuencias de los *Aotus* (Ao-DRB*W) y de las secuencias de humanos (HLA-DRB)



Promedio y Rango de homología entre Ao-DR β y HLA-DR β en el dominio β y posiciones de los Pockets

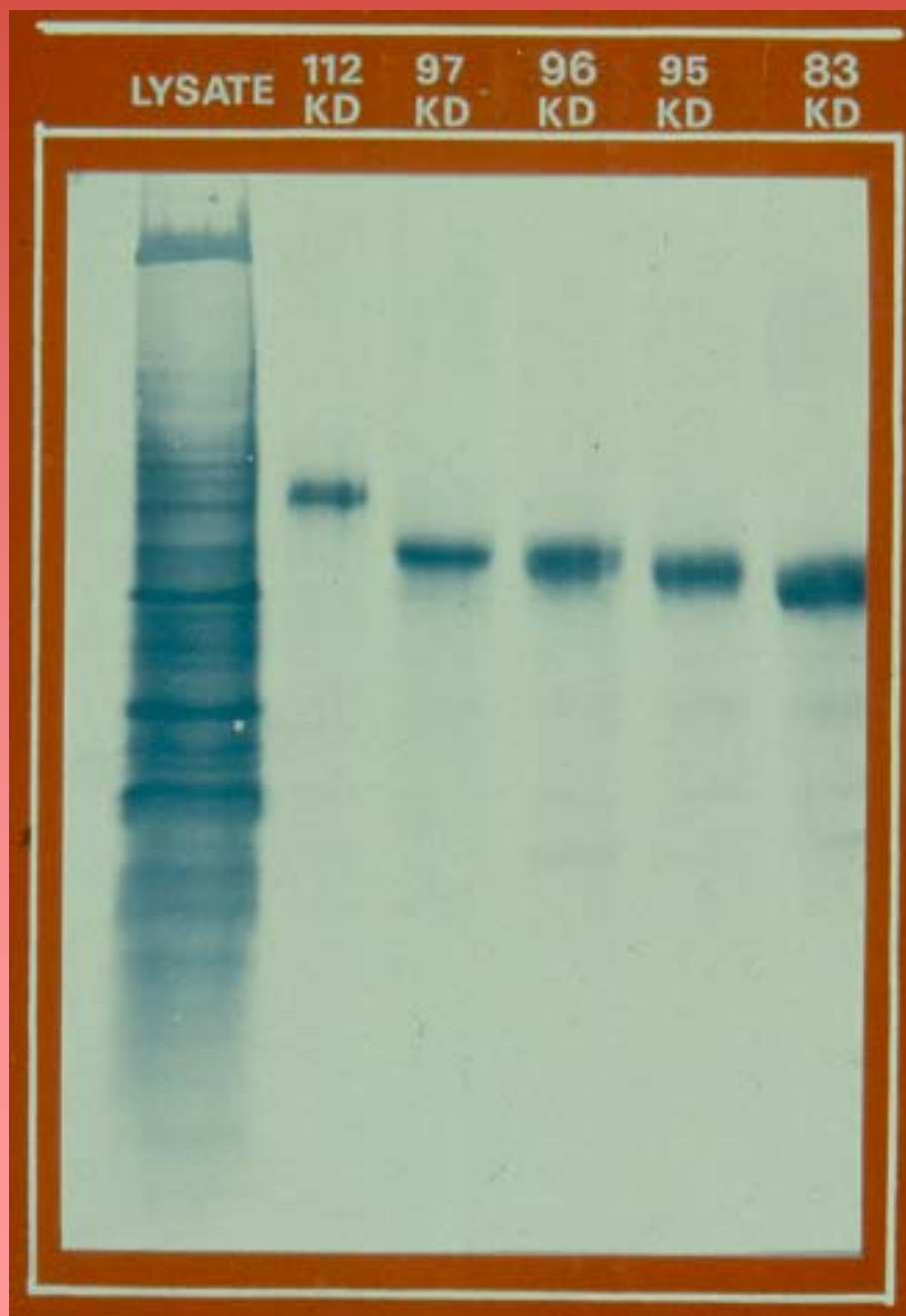
Owl Monkey Human		β domain			Pockets		
		% mean	% min	% max	% mean	% min	% max
Ao-DRB*W104	vs. HLA-DRB1*04	95	91	98	93	80	100
Ao-DRB*W304	vs. HLA-DRB1*03	92	88	95	85	77	91
Ao-DRB*W302	vs. HLA-DRB3	92	89	94	84	77	91
Ao-DRB*W11	vs. HLA-DRB1*08	92	88	96	83	74	91
Ao-DRB*W303	vs. HLA-DRB3	91	86	94	83	71	91
Ao-DRB*W301	vs. HLA-DRB3	90	86	93	84	74	91
Ao-DRB*W11	vs. HLA-DRB1*11	90	85	95	83	71	91
Ao-DRB*W11	vs. HLA-DRB1*13	90	85	96	81	71	91
Ao-DRB*W701/02	vs. HLA-DRB1*07	90	87	93	80	74	88
Ao-DRB*W101	vs. HLA-DRB1*01	90	84	94	80	74	88
Ao-DRB*W11	vs. HLA-DRB1*14	89	85	96	80	68	91
Ao-DRB*W4	vs. HLA DRB1*10	88	84	89	74	68	80
Ao-DRB*W4	vs. HLA DRB4	88	83	91	72	65	80





IMMUNOBLOT OF *P. falciparum* ISOLATED PROTEINS

Patarroyo M. E. et. al. 1987. *Vaccines*. 117-124



PARASITAEMIA IN MONKEYS IMMUNISED WITH MEROZOITE PURIFIED PROTEINS

Patarroyo M. E.. et al. 1987. *Vaccines* 87:117

Immunizing molecule	Monkey number	Percentage of parasitemia on days (after challenge)														
		6	7	8	9	10	11	12	13	14	15	16	17	18	19	
155K	087	0	0	0	0	0.88	0.30	2.0	3.0	5.35	8.8	Q				
	090	0	0	0	0	0	0	0.8	0.2	0.45	0.9	1.5	1.6	2.0	3.4	
115K	085	0	0.46	0.84	3.48	3.0	9.93	Q								
	150	0	0.05	0.20	1.12	1.67	nd	6.57	Q							
105K	165	0	0.33	1.92	3.72	10.9	Q									
	168	0	0.81	1.64	3.33	11.5	Q									
90K	160	0	0.13	1.2	3.39	4.14	13.65	Q								
	170	0	0.10	1.1	1.20	5.09	5.40	14.7	Q							
83K	125	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	130	0	0	0	0.6	0.2	0.3	3.63	1.6	1.1	0.6	nd	0	0	0	
	144	0	0	0	0	0	0	0	0.1	0.05	0.5	0.3	0.07	0.03	0.01	
60K	119	0	0	0	0.1	0.80	0.15	4.5	1.25	15.3	Q					
	122	0	0.13	0.25	0.15	0.80	2.5	17.5	Q							
55K	0.81	0	0	0	0	0	0.18	0.24	0.8	1.57	3.0	3.5	4.4	7.4	Q	
	102	0	0	0	0	0	0.05	0.05	0.5	0	0.1	0.6	0.9	5.4	7.3	
50K	0.29	0	0.40	1.50	3.6	4.67	10.0	Q								
	0.56	0	0.66	5.6	4.43	6.42	4.20	11.3	Q							
	131	0	0.31	1.50	0.50	3.93	2.0	16.5	Q							
40K	111	0	0	0	0	1.02	0.20	2.7	3.2	14.3	Q					
	114	0	0	0	0	0.8	0.5	2.0	2.0	2.0	7.6	Q				
35K	135	0	0	0	0	0	0	0.01	0.03	0.03	0.1	0.012	0.01	0	0	
	159	0	0.25	0.25	1.3	2.23	1.03	2.5	2.1	4.0	2.1	1.6	0.7	0.08	0.01	
30K	093	0	0.05	1.52	2.5	4.75	3.45	2.54	4.8	9.4	Q					
	171	0	0.1	2.5	1.6	2.63	2.05	13.0	Q							
23K	112	0	0.06	0	1.0	0	3.60	6.37	6.60	12.0	Q					
	137	0	0.05	0.05	0.20	0	1.1	3.2	nd	+						
Controls	501	0.40	0.76	3.0	2.0	7.82	9.07	Q								
	502	0.03	0.06	0.43	0.47	3.6	3.4	5.9	8.92	Q						
	199	1.0	4.3	3.11	9.8	26.0	Q									

nd indicates not determined

Handmade PEPTIDE SUPERMA

PEPTIDES

INSTRUCTIONS

WAVE COIN FOR SELECTION
OF FLAVORS. INSERT COIN &
WAIT FOR BEVERAGE. PLEASE
DO NOT PRESS A COIN INTO
THE MACHINE. THANK YOU.

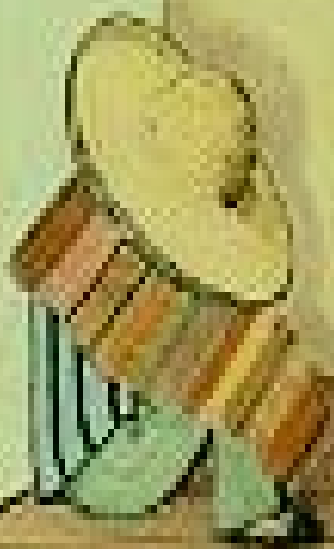
PLEASE
DO NOT
PRESS
COIN INTO
MACHINE

PEPTIDE
MACHINE

ALA	ARG	ASP	GLU	GLY
ILE	LEU	LYS	PHE	PRO
SER	THR	TRP	TYR	VAL
ASP	GLU	GLY	ILE	LEU
LYS	PHE	PRO	SER	THR
TRP	TYR	VAL	ALA	ARG
ASP	GLU	GLY	ILE	LEU
LYS	PHE	PRO	SER	THR
TRP	TYR	VAL	ALA	ARG

COIN
SLIT

PEPTIDE
MACHINE



Induction of protective immunity against experimental infection with malaria using synthetic peptides

**Manuel E. Patarroyo, Pedro Romero, Martha L. Torres,
Pedro Clavijo, Alberto Moreno, Alberto Martínez,
Raul Rodríguez, Fanny Guzman & Edelmira Cabezas**

**Instituto de Inmunología, Hospital San Juan de Dios,
Universidad Nacional de Colombia, Bogotá, Colombia**

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**Induction of protective immunity
against experimental infection
with malaria using synthetic peptides**

**Manuel E. Patarroyo, Pedro Romero, Martha L. Torres,
Pedro Clavijo, Alberto Moreno, Alberto Martínez,
Raul Rodríguez, Fanny Guzman & Edelmira Cabezas**

Instituto de Inmunología, Hospital San Juan de Dios,
Universidad Nacional de Colombia, Bogotá, Colombia

Synthetic peptides are potential vaccine candidates because they may be able to induce high antibody titres and specific cellular immune responses against native proteins and thus the whole invading organism¹. In a previous study we showed that immunization with molecules of relative molecular mass (M_r) 155,000 (155K) 83K, 55K and 35K, specific for the late schizont and merozoite stages of *Plasmodium falciparum*, could elicit either partial or total protection in *Aotus trivirgatus* monkeys experimentally infected with *P. falciparum*². Here we have chemically synthesized 18 peptides corresponding to different fragments of these proteins to immunize *Aotus trivirgatus* monkeys. Some peptides gave partial protection from challenge with *P. falciparum* parasites, but none provided complete protection individually. A combination of three partially protective peptides gave complete or almost complete protection, however, suggesting that this particular combination of peptides is a good candidate for a malaria vaccine.

AMINOACID SEQUENCE OF THE SYNTHETIC MALARIA VACCINE SPf66

Patarroyo M. et al. 1988. *Nature*. 332:158

S
 1 2 3 4 5 6 7 8 9 10 1 2 3 4 5
 Cys- Gly- Asp- Glu- Leu- Glu- Ala- Glu- Thr- Gln- Asn- Val - Tyr - Ala - Ala

 16 7 8 9 20 1 2 3 4 5 6 7 8 9 30
 Pro- Asn- Ala - Asn- Pro- Tyr - Ser - Leu- Phe- Gln - Lys- Glu- Lys- Met- Val

 1 2 3 4 5 6 7 8 9 40 1 2 3 4 5
 Leu - Pro- Asn- Ala - Asn- Pro- Pro- Ala - Asn- Lys- Lys- Asn- Ala - Gly- Cys
 S



**A synthetic vaccine protects humans
against challenge with asexual blood
stages of *Plasmodium falciparum* malaria**

**Manuel E. Patarroyo*, Roberto Amador*,
Pedro Clavijo*, Alberto Moreno*, Fanny Guzman*,
Pedro Romero*, Ricardo Tascon*, Antonio Franco*,
Luis A. Murillo*, Gabriel Ponton† & Gustavo Trujillo†**

* Instituto de Inmunologia, Hospital San Juan de Dios,
Universidad Nacional de Colombia, Bogota, Colombia SA.

† Hospital Militar Central de Colombia, Fuerzas Militares de
Colombia, Bogota, Colombia S.A.

We have previously shown that a mixture of three synthetic peptides (83.1, 55.1, 35.1), corresponding to fragments of the relative molecular mass 83,000 (83K), 55K and 35K *Plasmodium falciparum* merozoite-specific proteins, induces protection in *Aotus trivirgatus* monkeys experimentally infected with *P. falciparum*^{1,2}. Here we describe two polymeric synthetic hybrid proteins based on these peptides that delay or suppress the development of parasitaemia in immunized human volunteers.

Studies on the humoral immune response to a synthetic vaccine against *Plasmodium falciparum* malaria

M. SALCEDO, L. BARRETO, M. ROJAS, R. MOYA, J. COTE & M. E. PATARROYO

Institute of Immunology, Hospital San Juan de Dios, Universidad Nacional de Colombia, Bogotá, Colombia

(Accepted for publication 9 October 1990)

SUMMARY

A synthetic vaccine against the asexual blood stages of *P. falciparum*, the SPf 66 synthetic hybrid polymer, composed of peptides derived from three merozoite membrane proteins as well as one peptide from the sporozoite CS protein, has been developed by our group and tested in different protection assays in *Aotus* monkeys as well as in human volunteers. This study evaluates the humoral immune response induced by the SPf 66 protein vaccination in adult human volunteers from the Colombian Pacific coast as follows: determination of specific IgG antibody levels against SPf 66 by FAST-ELISA after each immunization; analysis of antibody reactivity with *P. falciparum* schizont lysates by immunoblots; and determination of the *in vitro* parasite growth inhibition. A clear boosting effect, dependent on time and dose, was observed in the antibody production kinetics. These antibodies also specifically recognize three proteins of the *P. falciparum* schizont lysate corresponding to the molecular weights of the proteins from which the amino acid sequence was derived. These sera were also capable of markedly inhibiting *in vitro* parasite growth.

Study of the safety and immunogenicity of the synthetic malaria SPf66 vaccine in children aged 1–14 years

Gloria Patarroyo, Lina Franco, Roberto Amador, Luis Angel Murillo, Claudia Lucia Rocha, Mauricio Rojas and Manuel Elkin Patarroyo*

Safety and immunogenicity tests of the SPf66 malaria vaccine have been carried out on a population of children, aged 1 to 14 years, in the town of Tumaco, Colombia. Adverse reactions measured after each vaccination were local and minimal, and observed in only a small percentage of the vaccinated children. One year later, no delayed reaction was evident. The majority of the child population developed high antibody titres against SPf66 and the degree of response did not vary with age. These induced antibodies recognize the native parasite proteins, in particular the molecules from which the amino acid sequence of this vaccine was deduced. These studies demonstrate that the SPf66 vaccine is safe and highly immunogenic for use in children >1 year old.

Safety and Immunogenicity of the Synthetic Malaria Vaccine SPf66 in a Large Field Trial

Roberto Amador, Alberto Moreno, Luis Angel Murillo,
Oscar Sierra, David Saavedra, Mauricio Rojas,
Ana Lucia Mora, Claudia Lucia Rocha,
Fernando Alvarado, Juan Carlos Falla,
Mauricio Orozco, Carlos Coronell, Norella Ortega,
Alberto Molano, José Fernando Velásquez,
María Victoria Valero, Lina Franco, Fanny Guzmán,
Luz Mary Salazar, Fabiola Espejo, Elsa Mora,
Rocío Farfán, Nohora Zapata, Jaiver Rosas,
Julio Cesar Calvo, Jaime Castro, Teódulo Quiñones,
Francisco Nuñez, and Manuel Elkin Patarroyo

*Instituto de Inmunología, Hospital San Juan de Dios,
Universidad Nacional de Colombia, Bogotá*

In the first field trial with synthetic malaria vaccine SPf66 in a large population naturally exposed to malaria, 9957 persons >1 year old and residing on the Colombian Pacific coast received three doses of the vaccine. To evaluate vaccine safety, clinical observations were made 30 min and 48 h after each immunization. There were no adverse reactions in 95.7% of cases. In the 4.3% of cases with adverse reactions, local induration and erythema were the most frequent. In a randomly selected group of vaccinees, anti-SPf66 antibody titers were measured after the third dose: 93% of the vaccinees raised antibodies to SPf66. Among these, 55% had titers >1:1600. These results demonstrate the safety and immunogenicity of the SPf66 vaccine in a large field trial.

THE LANCET

Vaccination with SPf66, a chemically synthesised vaccine, against *Plasmodium falciparum* malaria in Colombia

M. V. Valero L. R. Amador C. Galindo J. Figueroa
M. S. Bello L. A. Murillo A. L. Mora G. Patarroyo
C. L. Rocha M. Rojas J. J. Aponte L. E. Sarmiento
D. M. Lazada C. G. Coronell N. M. Ortega J. E. Rosas
P. L. Alonso M. E. Patarroyo

Reprinted from THE LANCET Saturday 20 March 1993

Preclinical and clinical studies have established the safety and immunogenicity of the chemically synthesised SPf66 malaria vaccine. The present study is a phase III randomised, double-blind, placebo-controlled, efficacy trial completed in La Tola, Colombia. 1548 volunteers over one year of age received three doses of either the vaccine (n = 738) or placebo (n = 810).

Active and passive case detection methods were used to document clinical episodes of malaria among the study population. The follow-up period began one month after the third dose and lasted for one year. 168 and 297 episodes of *Plasmodium falciparum* malaria were documented in the SPf66 group and the placebo group, respectively; this corresponds to a crude protective efficacy of 38·8%. Incidence rates for first or only *P. falciparum* malarial episodes were 22·3% per annum among the vaccine group and 44·3% among the placebo group. The efficacy against second episodes was 50·5% (95% CI 12·9–71·9). Therefore, the protective efficacy of SPf66 against first or only episodes of malaria was 38·8% (95% CI 12·9–71·9). The efficacy against second episodes was 50·5% (95% CI 12·9–71·9). The estimated protective efficacy against second episodes was 50·5% (95% CI 12·9–71·9).



A Population-Based Clinical Trial with the SPf66 Synthetic *Plasmodium falciparum* Malaria Vaccine in Venezuela

Oscar Noya G., Yleana Gabaldón Berti,
Belkisyolé Alarcón de Noya, Rafael Borges,
Noraida Zerpa, José David Urbáez, Alberto Madonna,
Enrique Garrido, M. Auxiliadora Jiménez,
Rafael E. Borges, Paul Garcia, Ivan Reyes,
Wolfgang Prieto, Cecilia Colmenares, Rosalba Pabón,
Tito Barraez, Lucía G. de Caceres, Nabor Godoy,
and Rolando Sifontes

Instituto de Medicina Tropical and Escuela de Salud Pública,
Universidad Central de Venezuela, Caracas; Dirección de Endemias
Rurales, Malariología, Ministerio de Sanidad y Asistencia Social,
Maracay, Venezuela

A phase III malaria vaccine trial in 13 villages in an endemic area, South Venezuela, compared incidence rates of *Plasmodium falciparum* and *Plasmodium vivax* infections in 1422 vaccinated and 938 nonvaccinated subjects over 18 months. The SPf66 vaccine was given in three doses, on days 0, 20, and 112. Vaccination was complete in 976 subjects (68.7%). Minor side effects requiring no treatment were reported by 123 (12.6%), with an apparent increase in frequency from the first to the third vaccine dose. No autoimmune evidence was observed in a sample of subjects. Antibodies against SPf66 were present at low titers in 24.7% of tested subjects before vaccination, increasing to 53.6% after the second dose and to 73.6% after the third dose; 26.4% of subjects initially seronegative never seroconverted. The SPf66 malaria vaccine showed a protective efficacy of 55% (95% confidence interval, 21%–75%) against *P. falciparum* and of 41% (19%–57%) against *P. vivax* malaria.

Safety, immunogenicity and protective effect of the SPf66 malaria synthetic vaccine against *Plasmodium falciparum* infection in a randomized double-blind placebo-controlled field trial in an endemic area of Ecuador

Fernando Sempértegui^{*†}, Bertha Estrella^{*}, Juan Moscoso^{*}, Luis Piedrahita C.^{*}, Denise Hernández^{*}, José Gaybor^{*}, Plutarco Naranjo^{*}, Olmedo Mancero^{*}, Silvio Arias^{*}, Rodolfo Bernal^{*}, Maria E. Córdova^{*}, Jorge Suárez^{*} and Fabio Zicker[†]

A total of 537 subjects were randomized to receive either SPf66 malaria vaccine against Plasmodium falciparum or placebo in three doses (days 0, 30 and 180). Subjects completing the course of vaccination (230 in the vaccine and 238 in the placebo group) were followed up for a further 12 months. Case detection surveillance was implemented by parasitological cross-sectional surveys every 2 months and by monthly household visits to each participant. Symptomatic subjects were also diagnosed in a local health centre. Minor local side-effects were observed mainly after the second dose in about 19% of the vaccinated subjects and in 3.7% of the placebo group. Thirty days after the third dose the prevalence of anti-SPf66 antibodies was 57% in the vaccine and 8.8% in the placebo groups. The prevaccination prevalence of antibodies measured by indirect immunofluorescence assay increased with age and seemed to be inversely related to anti-SPf66 antibody production. Immune response to SPf66 was independent of age. Vaccine efficacy was calculated based on person-time of exposure. The protective effect considering any malaria episode was 66.8% (confidence interval = -2.7-89.3%) and considering only one episode per individual was 60.2% (95% confidence interval = -26-87.5%).





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Evaluation of SPf66 malaria vaccine during a 22-month follow-up field trial in the Pacific coast of Colombia

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A double-blind randomized placebo-controlled field trial with the SPf66 malaria vaccine was carried out in an endemic area consisting of 14 small villages with exclusive fluvial access, in a rain forest area along the Rosario River, Colombia. A total of 1257 subjects completed the full three dose vaccination schedule on days 0, 30 and 180 (643 vaccinated group/623 placebo group) and were followed-up by passive and active surveillance over a period of 22 months. One hundred and thirty-four Plasmodium falciparum malaria episodes were detected (53 in vaccinated group/81 in placebo group), yielding an attack rate of 5.47 cases/100 person years of follow-up (pyears) in the vaccine group and 8.44/100 pyears in the placebo group. The estimated vaccine protective efficacy was 35.2% (95% CI 8.4–54.2%, $P=0.01$). This result supports earlier findings that the SPf66 malaria vaccine diminishes the risk of infection by P. falciparum in endemic areas of South America. Copyright © 1996 Elsevier Science Ltd.



Randomised trial of efficacy of SPf66 vaccine against *Plasmodium falciparum* malaria in children in southern Tanzania

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Summary

Effective, safe antimalarial vaccines have proved elusive. The synthetic polypeptide SPf66 vaccine is based on pre-erythrocytic and asexual blood-stage proteins of *Plasmodium falciparum*. We report here a randomised double-blind placebo-controlled trial of the efficacy of the SPf66 vaccine against clinical *P falciparum* malaria in Idete, southern Tanzania, an area of intense perennial malaria transmission.

586 children aged 1-5 years received three doses of vaccine (n=274) or placebo (n=312). The incidence and density of parasitaemia were assessed through repeated cross-sectional surveys on subgroups of children. Morbidity was monitored over a 1 year period through passive case detection in all children plus active case detection in a subgroup of 191. An episode of clinical malaria was defined as measured fever ($\geq 37.5^{\circ}\text{C}$) and parasite density $>20\,000/\mu\text{L}$.

No severe side-effects were seen and the frequency of mild side-effects after the third dose was less than 6%. The vaccine was highly immunogenic and after three doses all vaccine recipients had detectable anti-SPf66 antibodies: the geometric mean index of response was 8.3 in the vaccine group and 0.7 in the placebo group. The incidence of parasitaemia was similar in both groups. 123 children had at least one episode of clinical malaria during the follow-up period after the third dose and annual incidence rates were 0.25 in the vaccine group and 0.35 in the placebo group. Estimated vaccine efficacy was 31% (95% confidence interval 0-52%; $p=0.046$). After the third dose there were 6 deaths among the study cohort (1 vaccine, 5 placebo). This study confirms that SPf66 is safe, immunogenic and reduces the risk of clinical malaria among children exposed to intense *P falciparum* transmission.

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Duration of Protection and Age-Dependence of the Effects of the SPf66 Malaria Vaccine in African Children Exposed to Intense Transmission of *Plasmodium falciparum*

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The SPf66 synthetic vaccine is safe and partly efficacious against *Plasmodium falciparum* malaria among children 1–5 years old. The estimated vaccine efficacy [VE] for all clinical episodes over a period of 18 months after the third dose is 25% (95% confidence interval [CI], 1%–44%; $P = .044$). The observed temporal variations in efficacy could have been due to chance (likelihood ratio $\chi^2 = 13.8$, 8 df; $P = .086$). Efficacy against clinical malaria did not vary significantly with age ($\chi^2 = 1.07$, 4 df; $P = .90$). Overall parasite density was 21% lower in vaccine recipients than in the placebo group (95% CI, 0%–38%; $P = .044$). Further development of SPf66 may require trials to evaluate safety, immunogenicity, and efficacy when administered in the first year of life, together with other vaccines contained in the Expanded Programme of Immunization schedule.

Analysis of Multiple *Plasmodium falciparum* Infections in Tanzanian Children during the Phase III Trial of the Malaria Vaccine SPf66

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In the first phase III efficacy trial of the malaria vaccine SPf66 in Africa, MOIs in SPf66- and placebo-vaccinated children were analyzed by polymerase chain reaction–restriction fragment length polymorphism of the *Plasmodium falciparum* merozoite surface antigen 2 (MSA2). MOIs were significantly reduced in asymptomatic vaccine recipients compared with those in asymptomatic placebo recipients; however, no differences were observed among symptomatic children in the vaccine and control groups. These results show that immunization with SPf66 modulates the course of naturally occurring infections, as reflected by reduced MOIs. In placebo recipients, however, there was a significant negative correlation between numbers of infecting genotypes, as identified by MSA2, and morbidity. Asymptomatic placebo recipients had an average of 5 concurrent infections, whereas children with clinical cases had an average of 3.4 infections. These data provide further evidence that premunition from concurrent infections is important in immunity against clinical malaria. No such effect of multiple infections was found in the vaccinated group.

Randomised double-blind placebo-controlled trial of SPf66 malaria vaccine in children in northwestern Thailand

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Summary

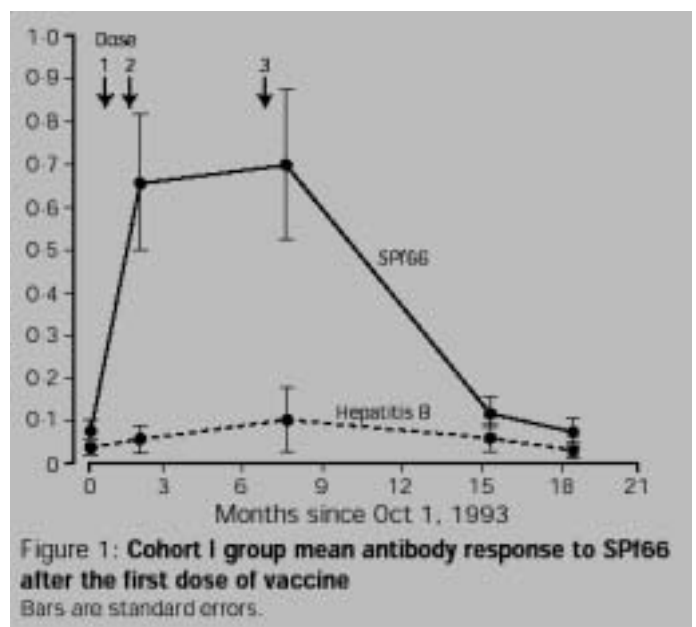
Background Previous efficacy trials of SPf66 malaria vaccine have produced conflicting results in different populations. We report a randomised double-blind trial of the SPf66 vaccine conducted in Karen children aged 2–15 living in a malarious region of northwestern Thailand. Recombinant hepatitis B vaccine was used as a comparator.

Methods The study had a power of 90% to detect an efficacy of 30%, defined as a reduction in the incidence of first cases of symptomatic falciparum malaria after three doses of vaccine. 1221 children received three immunisations and were eligible for the primary efficacy analysis. Intense active and passive case detection continued over 15 months of follow-up.

Findings The SPf66 vaccine was well tolerated, although 26 children had mild or moderately severe local or systemic allergic reactions, compared with none in the comparator group. The vaccine was immunogenic; after three doses, 73% of recipients had seroconverted. There were no deaths due to malaria during the study. During the 15-month period of evaluation there were 379 first cases of symptomatic falciparum malaria (195 in the SPf66 recipients, 184 in the comparator group); an SPf66 efficacy of –9% (95% CI –33 to 14, $p=0.41$). No significant differences between the two study groups in parasite density or any other measure of malaria-related morbidity were detected.

Interpretation These findings are consistent with a recent study showing lack of efficacy of SPf66 among Gambian infants and differ from earlier positive reports from South America and evidence of borderline efficacy from Tanzania.

We conclude that SPf66 does not protect against clinical falciparum malaria and that further efficacy trials are not warranted.



The vaccine lots used in this study were produced and formulated in the USA, whereas the previous studies used vaccine produced in Colombia. At the request of the WHO, samples from both sources have undergone extensive independent characterisation by the National Institute of Biological Standards and Controls, UK, and remarkable consistency has been found between lots of bulk and formulated vaccine produced by the two laboratories.

In our hands, the ratio of monomer to polymer was found to be important in the context of peptide binding to alum and thus to immunogenicity. Our SPf66 formulations were prepared with a higher alum concentration per dose than used in Colombia to achieve optimum peptide binding ratios (>50%). The seroconversion rates with standard

A pilot safety and immunogenicity study of the malaria vaccine SPf66 in Gambian infants

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SUMMARY

A pilot safety and immunogenicity trial of the malaria vaccine SPf66 has been undertaken in 150 Gambian infants. No significant systemic side effects were recorded but modest local reactions were seen after the administration of a third 1.0 mg dose. SPf66 produced in Colombia was more immunogenic than SPf66 produced in the USA and a 1.0 mg dose of each vaccine gave higher antibody levels than a 0.5 mg dose. However, antibody levels fell rapidly after administration of the third dose of vaccine and showed little change over the following malaria transmission season.

Comment

The SPf66 Malaria Vaccine: What is the Evidence for Efficacy?

P. Graves, H. Gelband and P. Garner

Results of individual trials were combined using the Peto odds ratio (OR), for which the trials are weighted by the inverse of their variance⁶. The weighting thus takes into account the number of participants in a trial and the number of outcomes observed. The OR result may be interpreted as the relative risk in the vaccinated group compared with the control group. An OR significantly less than 1.0 (shown as the vertical line in Table 1) indicates a protective effect of the vaccine. The OR is converted to vaccine efficacy (%) by the following formula: Efficacy = $(1 - \text{OR}) \times 100$.

Parasitology Today, vol. 14, no. 6, 1998

Systematic review findings. In the current edition of the Cochrane malaria vaccine review, six trials of SPf66 met the inclusion criteria: three from South America⁷⁻⁹, two from Africa^{10,11} and one from Thailand³ (Table 2). Two other trials in South America were excluded due to lack of proper randomization. Results are presented in Table 1 (upper part) for the incidence of the first or only attack of clinical malaria. The Peto OR was 0.77 [95% confidence interval (CI) = 0.68–

0.88], giving a combined estimate of vaccine efficacy of 23% (95% CI = 12–32%).

During the follow-up period, some participants had more than one

episode of clinical malaria, and five of the six SPf66 trials reported the total number of episodes. The results from these five trials suggest that vaccine efficacy was greater against second or subsequent clinical episodes than it was in preventing malaria completely. The overall (unweighted) incidence of the first or only episode of malaria was reduced from 240 per 1000 in the placebo groups to 191 per 1000 in the

SPf66 groups (a reduction of 20%), whereas the incidence of subsequent attacks was reduced from 62 per 1000 to 41 per 1000 (a reduction of 34%).

Only two trials reported on the most

Reviews

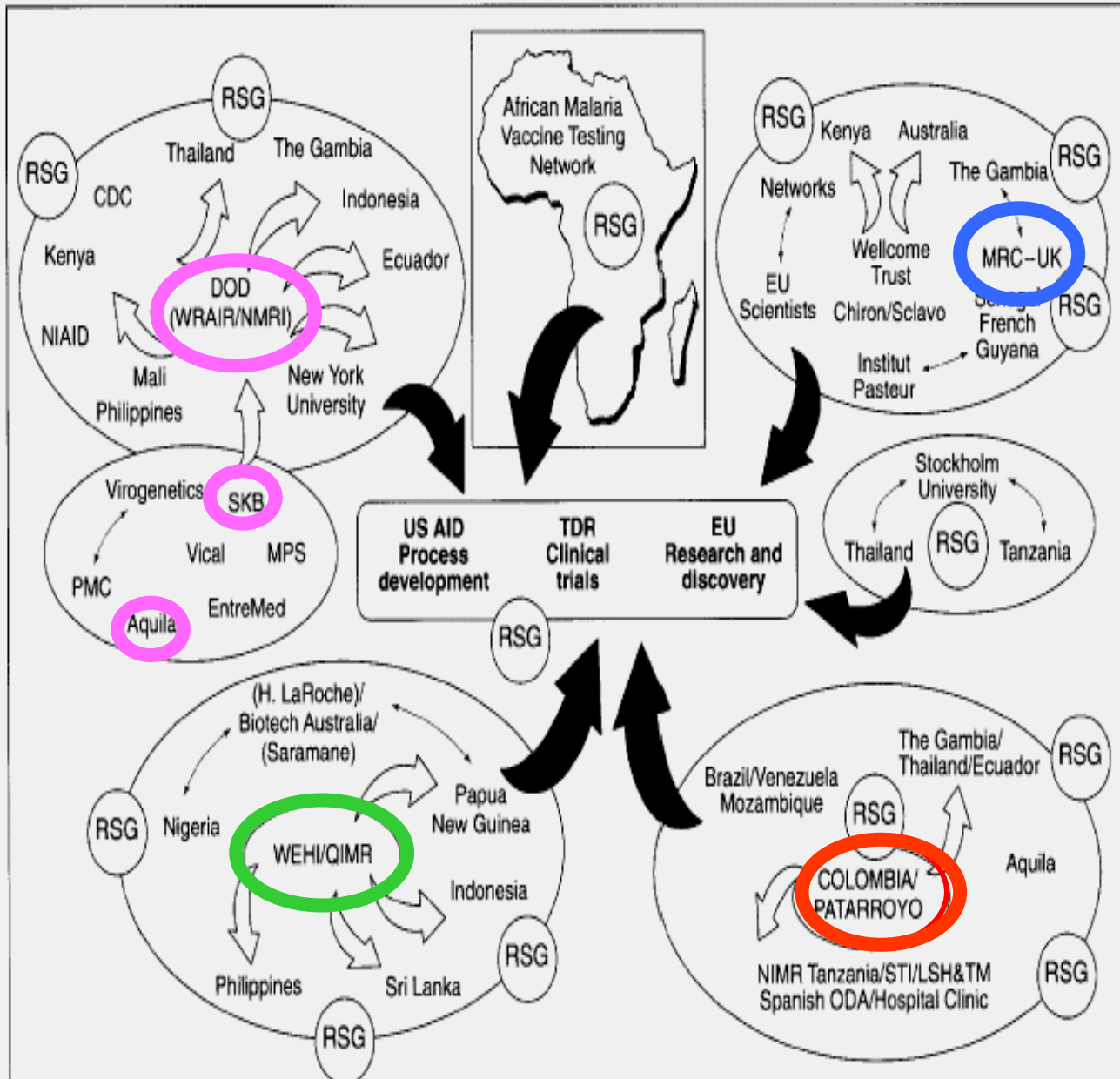


Fig. 2. A global strategy for malaria vaccine development. This schema illustrates the major contributors to malaria vaccine development. Abbreviations: CDC, Centers for Disease Control; DOD, Dept of Defence; EU, European Union; LSH&TM, London School of Hygiene and Tropical Medicine; MPS, Mimotope Peptide Systems; MRC, Medical Research Council; NIAID, National Institutes of Allergy and Infectious Diseases; NIMR, National Institute for Medical Research, Tanzania; NMRI, Naval Malaria Research Institute; ODA, Overseas Development Agency; PMC, Pasteur/Merieux/Connaught; QIMR, Queensland Institute for Medical Research; RSG, Research Strengthening Group of TDR; SKB, SmithKline Biologicals; STI, Swiss Tropical Institute; TRAP, thrombospondin-related anonymous protein; WEHI, Walter and Elisa Hall Institute; WRAIR, Walter Reed Army Institute for Research.

Papers

Safety, immunogenicity and limited efficacy study of a recombinant *Plasmodium falciparum* circumsporozoite vaccine in Thai soldiers

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Thai soldiers were vaccinated with a recombinant protein derived from the central repeat region of the circumsporozoite (CS) protein of *Plasmodium falciparum* conjugated to Toxin A (detoxified) of *Pseudomonas aeruginosa* (R32Tox-A) to evaluate its safety, immunogenicity and efficacy. In a randomized, double-blind manner, 199 volunteers received either R32Tox-A or a control vaccine at 0, 8, and 16 weeks. Immunization was performed in a malaria non-transmission area, after completion of which volunteers were deployed to an endemic border area and monitored closely to allow early detection and treatment of infection. The vaccine was found to be safe and to elicit antibody responses in all vaccinees. Peak CS antibody (IgG) concentrations in malaria-experienced vaccinees exceeded those in malaria-naïve vaccinees (mean 40.6 versus 16.1 $\mu\text{g ml}^{-1}$; $p = 0.005$) as well as those induced by previous CS protein-derived vaccines and observed in association with natural infections. A low number of malaria cases occurred during the study. Serum malaria revealed no differences between vaccinated and non-vaccinated subjects. Secondary analyses revealed that CS antibody levels were lower in vaccinee malaria cases than in non-cases, 3 and 5 months after the third dose of vaccine ($p = 0.06$ and $p = 0.014$, respectively). Because antibody levels had fallen substantially before peak malaria transmission occurred, the question of whether high levels of CS antibody are protective remains to be resolved.

***Plasmodium falciparum* circumsporozoite vaccine immunogenicity and efficacy trial with natural challenge quantitation in an area of endemic human malaria of Kenya**

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J.C. Sadoff***

It has been hypothesized that antibody induced by Plasmodium falciparum circumsporozoite protein vaccine would be effective against endemic human malaria. In a malaria endemic region of Kenya, 76 volunteers, in 38 pairs sleeping adjacently, were immunized with subunit circumsporozoite protein Asn-Ala-Asn-Pro tetrapeptide repeat-pseudomonas toxin A, or hepatitis B vaccine. After quinine and doxycycline, volunteers were followed for illness daily, parasitemia weekly, antibody, T-lymphocyte responses, and treated if indicated. Anopheles mosquitoes resting in houses were collected, and tested for P. falciparum antigen, or dissected for sporozoites and tested for blood meal ABO type and P. falciparum antigen. Vaccine was safe, with side-effects similar in both groups, and immunogenic, engendering IgG antibody as high as $600 \mu\text{g ml}^{-1}$, but did not increase the proportion of volunteers with T-lymphocyte responses. Estimation of P. falciparum challenge averaged 0.194 potentially infective Anopheles bites/volunteer/day. Mosquito blood meals showed no difference in biting intensity between vaccine and control groups. Both groups had similar malaria-free survival curves, cumulative positive blood slides, cumulative parasites mm^{-3} , and numbers of parasites mm^{-3} on first positive blood slide, during three post-vaccination observation periods. Every volunteer had P. falciparum parasitemia at least once. Vaccinees had 82% and controls 89% incidences of symptomatic parasitemia ($P=0.514$, efficacy 9%, statistical power 95% probability of efficacy $<50\%$). Vaccine-induced anti-sporozoite antibody was not protective in this study. Within designed statistical precisions the present study is in agreement with efficacy studies in Colombia, Venezuela and Tanzania.

Phase I/IIa Safety, Immunogenicity, and Efficacy Trial of NYVAC-Pf7, a Pox-Vectored, Multiantigen, Multistage Vaccine Candidate for *Plasmodium falciparum* Malaria

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Candidate malaria vaccines have failed to elicit consistently protective immune responses against challenge with *Plasmodium falciparum*. NYVAC-Pf7, a highly attenuated vaccinia virus with 7 *P. falciparum* genes inserted into its genome, was tested in a phase I/IIa safety, immunogenicity, and efficacy vaccine trial in human volunteers. Malaria genes inserted into the NYVAC genome encoded proteins from all stages of the parasite's life cycle. Volunteers received three immunizations of two different dosages of NYVAC-Pf7. The vaccine was safe and well tolerated but variably immunogenic. While antibody responses were generally poor, cellular immune responses were detected in >90% of the volunteers. Of the 35 volunteers challenged with the bite of 5 *P. falciparum*-infected *Anopheles* mosquitoes, 1 was completely protected, and there was a significant delay in time to parasite patency in the groups of volunteers who received either the low or high dose of vaccine compared with control volunteers.

Effect of vaccination with 3 recombinant asexual-stage malaria antigens on initial growth rates of *Plasmodium falciparum* in non-immune volunteers

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Abstract

A placebo controlled, randomised, double blind trial was conducted in human volunteers to test a mixture of three recombinant *Plasmodium falciparum* blood stage antigens for its ability to reduce the initial growth rates of parasites. The vaccine contained recombinant MSP2 (3D7 allele), a portion of MSP1 (190LCS.T3) and part of the RESA antigen (C terminal 771 amino acids) in the Montanide ISA 720 adjuvant (SEPPIC). Twelve volunteers received two doses of the vaccine, 6 weeks apart. The five participants in the placebo group received an equivalent volume of the adjuvant emulsion using the same schedule. Antibody responses were low, as has been reported in earlier studies with this combination, while T cell responses were stronger. All the volunteers were challenged with approximately 140 ring infected red cells of the 3D7 cloned line, 4 weeks after the second dose. Parasitaemia was determined once daily from day 4 using a sensitive and quantitative PCR assay. All the volunteers were infected and were treated on day 8, before any developed symptoms. There was no significant difference in initial parasite growth rates between the verum and placebo groups, nor was there any significant correlation between parasite growth rates and any of the measured immunological responses. These results suggest that the formulation tested in this trial did not generate immune responses that were strong enough to reduce parasite growth in naive volunteers.

Enhanced T-cell immunogenicity of plasmid DNA vaccines boosted by recombinant modified vaccinia virus Ankara in humans

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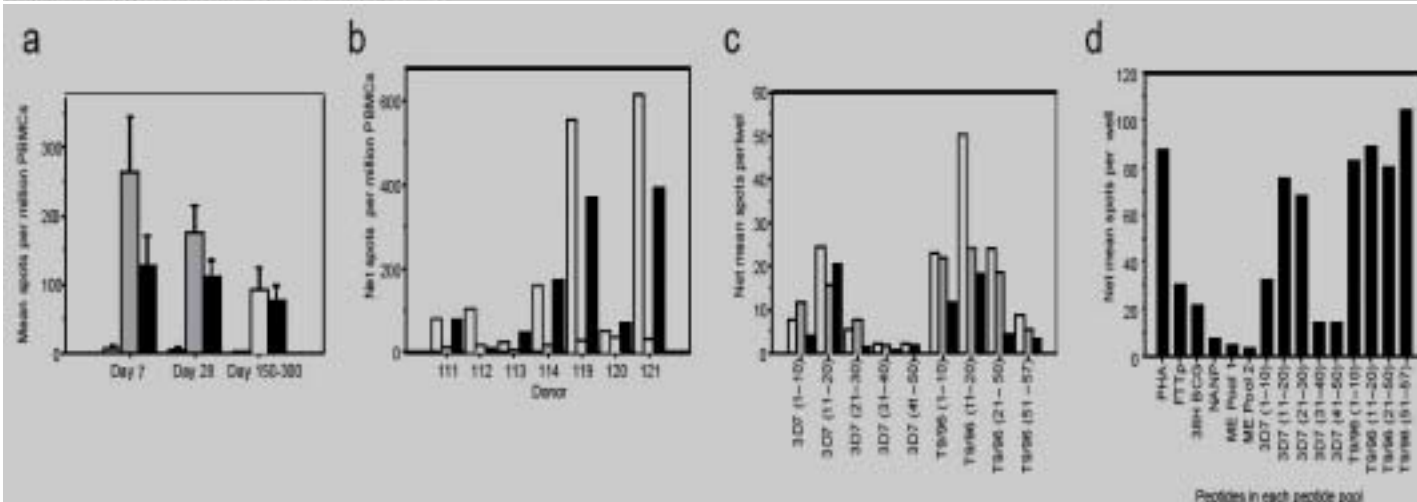
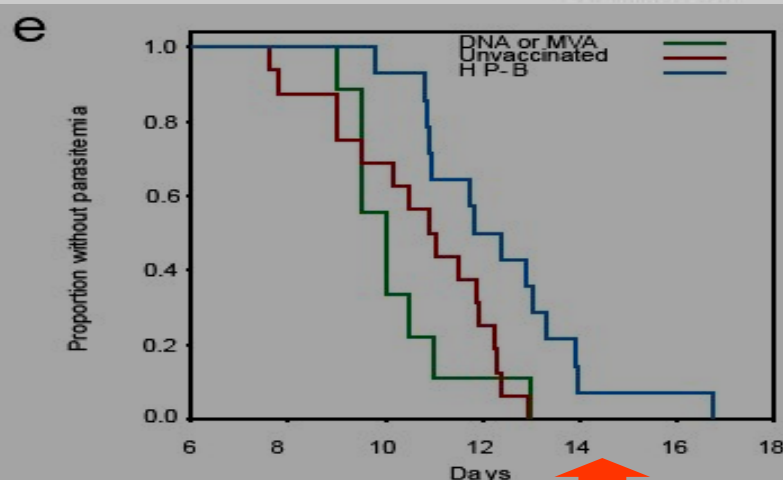


Figure 3 Characteristics of induced T-cell responses and protective efficacy. (a) Time course at 7, 28 and 150–300 d after vaccination for six subjects in group GGMM33. □, ME epitopes; ▨, TRAP TR96 peptides; ■, TRAP 3D7 peptides. The arithmetic mean and standard error are shown. (b) The results of CD4⁺ and CD8⁺ T-cell depletion experiments from GDD/MM1/5 and GDDMM3/5 groups 7 d after the last vaccine. □, undepleted; ▨, CD4-depleted; ■, CD8-depleted. (c,d) Breadth of the induced responses. The responses to each peptide pool shown in c are from the same subjects and time points as in a. □, day 7; ▨, day 28; ■, days 150–300. Data in d are 7 d after the first MVA booster dose of 1.5×10^7 PFU, for those who had these priming doses of 2 mg of DNA/ME-TRAP intramuscularly. The nomenclature of the peptide pools is described in Fig. 2b and in the footnote to Table 1. (e) Kaplan–Meier curves of time from sporadic challenge to parasitemia detected on thick blood film for three groups: 16 unvaccinated control subjects, 14 heterologous prime-boost vaccinated subjects who received either GGMM33 ($n=6$), GDDMM3/5 ($n=4$) or GDD/MM1/5 ($n=4$), these being the groups with the strongest immune responses, and 9 vaccinated subjects who received just one of the vaccines, either MM33 ($n=4$) or GDD1/5 ($n=5$; Table 1). HP-B, heterologous prime-boost. Development of parasitemia is delayed significantly in the heterologous prime-boost immunization group (log rank test, $P=0.013$) compared with the unvaccinated controls, but there is no significant difference in time to parasitemia when comparing the volunteers who received DNA only or MVA only and the unvaccinated group.



Safety, immunogenicity and efficacy of a pre-erythrocytic malaria candidate vaccine, ICC-1132 formulated in Seppic ISA 720

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Abstract

ICC-1132, a recombinant protein comprising a modified hepatitis B core protein with a B cell (NANP) and two T cell epitopes of *Plasmodium falciparum* circumsporozoite protein (CSP), was administered i.m. as a single 50 µg dose in Seppic ISA 720 to 11 volunteers. Local reactogenicity and systemic side effects were acceptable with the predominant finding being mild pain at the injection site. This regimen induced anti-NANP antibodies in 10/11 and modest T cell response. There was no evidence of protection from experimental challenge with *P. falciparum* sporozoites. Other formulations and/or multi-dose regimens will be required to enhance the immunogenicity and efficacy of ICC-1132.

Efficacy of the RTS,S/AS02A vaccine against *Plasmodium falciparum* infection and disease in young African children: randomised controlled trial

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Summary

Background Development of an effective malaria vaccine could greatly contribute to disease control. RTS,S/AS02A is a pre-erythrocytic vaccine candidate based on *Plasmodium falciparum* circumsporozoite surface antigen. We aimed to assess vaccine efficacy, immunogenicity, and safety in young African children.

Methods We did a double-blind, phase IIb, randomised controlled trial in Mozambique in 2002 children aged 1–4 years. The study included two cohorts of children living in two separate areas which underwent different follow-up schemes. Participants were randomly allocated three doses of either RTS,S/AS02A candidate malaria vaccine or control vaccines. The primary endpoint, determined in cohort 1 (n=1605), was time to first clinical episode of *P falciparum* malaria (axillary temperature $\geq 37.5^{\circ}\text{C}$ and *P falciparum* asexual parasitaemia >2500 per μL) over a 6-month surveillance period. Efficacy for prevention of new infections was determined in cohort 2 (n=417). Analysis was per protocol.

Findings 115 children in cohort 1 and 50 in cohort 2 did not receive all three doses and were excluded from the per-protocol analysis. Vaccine efficacy for the first clinical episodes was 29.9% (95% CI 11.0–44.8; $p=0.004$). At the end of the 6-month observation period, prevalence of *P falciparum* infection was 37% lower in the RTS,S/AS02A group compared with the control group (11.9% vs 18.9%; $p=0.0003$). Vaccine efficacy for severe malaria was 57.7% (95% CI 16.2–80.6; $p=0.019$). In cohort 2, vaccine efficacy for extending time to first infection was 45.0% (31.4–55.9; $p<0.0001$).

Interpretation The RTS,S/AS02A vaccine was safe, well tolerated, and immunogenic. Our results show development of an effective vaccine against malaria is feasible.

Malaria vaccine: 3 or 6 months' protection?

The results of the trial of the RTS,S/AS02A malaria vaccine reported by Alonso and colleagues (Oct 16, p 1411)¹ are exciting and show clearly a protective effect against severe malaria and delays in the time to a first attack of clinical malaria and to the appearance of parasites in the blood after vaccination. However, the authors' conclusions that protection against clinical disease showed no evidence of waning during the 6 month follow-up and that there was only limited waning of the efficacy against malaria infection during this period should be interpreted with caution, particularly since an earlier trial² with this vaccine in adults in The Gambia showed evidence of a protective effect against malaria infection that lasted for only 2–3 months. Although the new data are compatible with sustained protec-

tion over 6 months, they are also compatible with there being little protection against clinical malaria or malaria infection for more than 3 months after vaccination. The data in figure 4 in the report, for example, show that among those who had not developed clinical disease by 3 months after vaccination, the risk of an episode of malaria in the next 3 months was similar (about 7–8%) in vaccinated and unvaccinated groups. Also, the risk of acquiring infection by 6 months among those not infected at 3 months was similar in both groups (about 65%) and, overall, during the 6-month period similar numbers of children were infected (157 and 166, respectively).

Additionally, there is little evidence of protection against clinical episodes of malaria other than the first. After a first episode the rate of subsequent

episodes was slightly higher in the vaccinated group than in the unvaccinated group (deduced from table 3: 30 episodes in 19.4 person years at risk [PYAR; rate 1.5 per year] in the vaccinated group compared with 31 episodes in 27.2 PYAR [rate 1.1 per year] in the unvaccinated group).

*P G Smith, P J Milligan
Peter.smith@lshtm.ac.uk

MRC Tropical Epidemiology Group, London School of Hygiene and Tropical Medicine, London WC1E 7HT, UK

- 1 Alonso PL, Sacarlal J, Aponte JJ, et al. Efficacy of the RTS,S/AS02A vaccine against *Plasmodium falciparum* infection and disease in young African children: randomised controlled trial. *Lancet* 2004; **364**: 1411–20.
- 2 Bojang KA, Milligan PJ, Pinder M, et al. Efficacy of RTS,S/AS02 malaria vaccine against *Plasmodium falciparum* infection in semi-immune adult men in The Gambia: a randomised trial. *Lancet* 2001; **358**: 927–34.

Efficacy of RTS,S/AS02 malaria vaccine against *Plasmodium falciparum* infection in semi-immune adult men in The Gambia: a randomised trial

Kalifa A Bojang, Paul J M Milligan, Margaret Pinder, Laurence Vigneron, Ali Allouche, Kent E Kesteven, **Ripley Ballou**, David J Conway, William H H Reece, Philip Gothard, Lawrence Yamuah, Martine Delchambre, Gerald Voss, Brian M Greenwood, Adrian Hill, Keith P W J McAdam, Nadia Tornieporth, Joe D Cohen, and Tom Doherty, for the RTS,S Malaria Vaccine Trial Team*

Findings 250 men (131 in the RTS,S/AS02 group and 119 in the control group) received three doses of vaccine and were followed up for 15 weeks. RTS,S/AS02 was safe and well tolerated. *P falciparum* infections occurred significantly earlier in the control group than the RTS,S/AS02 group (Wilcoxon's test $p=0.018$). Vaccine efficacy, adjusted for confounders, was 34% (95% CI 8.0–53, $p=0.014$). Protection seemed to wane: estimated efficacy during the first 9 weeks of follow-up was 71% (46–85), but decreased to 0% (–52 to 34) in the last 6 weeks. Vaccination induced strong antibody responses to circumsporozoite protein and strong T-cell responses. Protection was not limited to the NF54 parasite genotype from which the vaccine was derived. 158 men received a fourth dose the next year and were followed up for 9 weeks; during this time, vaccine efficacy was 47% (4–71, $p=0.037$).

Comment

The SPf66 Malaria Vaccine: What is the Evidence for Efficacy?

P. Graves, H. Gelband and P. Garner

Results of individual trials were combined using the Peto odds ratio (OR), for which the trials are weighted by the inverse of their variance⁶. The weighting thus takes into account the number of participants in a trial and the number of outcomes observed. The OR result may be interpreted as the relative risk in the vaccinated group compared with the control group. An OR significantly less than 1.0 (shown as the vertical line in Table 1) indicates a protective effect of the vaccine. The OR is converted to vaccine efficacy (%) by the following formula: Efficacy = $(1 - \text{OR}) \times 100$.

Parasitology Today, vol. 14, no. 6, 1998

Systematic review findings. In the current edition of the Cochrane malaria vaccine review, six trials of SPf66 met the inclusion criteria: three from South America⁷⁻⁹, two from Africa^{10,11} and one from Thailand³ (Table 2). Two other trials in South America were excluded due to lack of proper randomization. Results are presented in Table 1 (upper part) for the incidence of the first or only attack of clinical malaria. The Peto OR was 0.77 [95% confidence interval (CI) = 0.68–

0.88], giving a combined estimate of vaccine efficacy of 23% (95% CI = 12–32%).

During the follow-up period, some participants had more than one

episode of clinical malaria, and five of the six SPf66 trials reported the total number of episodes. The results from these five trials suggest that vaccine efficacy was greater against second or subsequent clinical episodes than it was in preventing malaria completely. The overall (unweighted) incidence of the first or only episode of malaria was reduced from 240 per 1000 in the placebo groups to 191 per 1000 in the

SPf66 groups (a reduction of 20%), whereas the incidence of subsequent attacks was reduced from 62 per 1000 to 41 per 1000 (a reduction of 34%).

Only two trials reported on the most

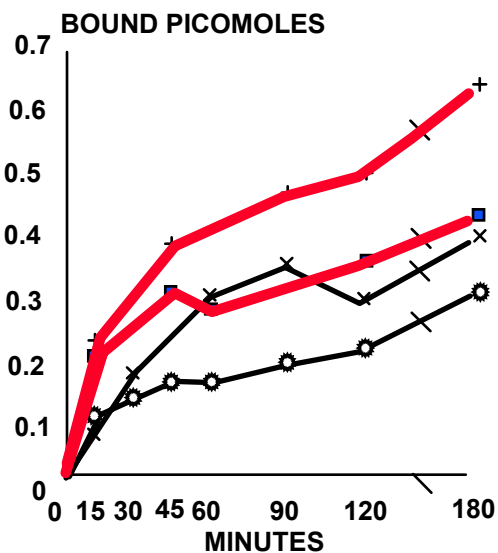
COLFAVAC

The COLombian FAlciparum VACcine

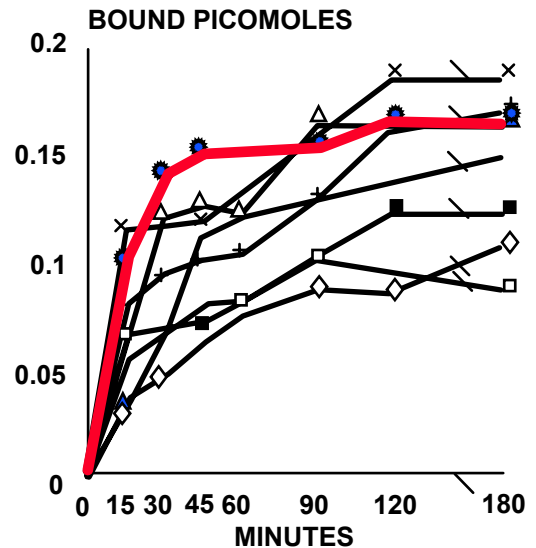
SPECIFIC UPTAKE OF THE DIFFERENT PEPTIDES BY HUMAN R.B.C.

Calvo M. et al. Peptide Research, 1991, 4, 324-333

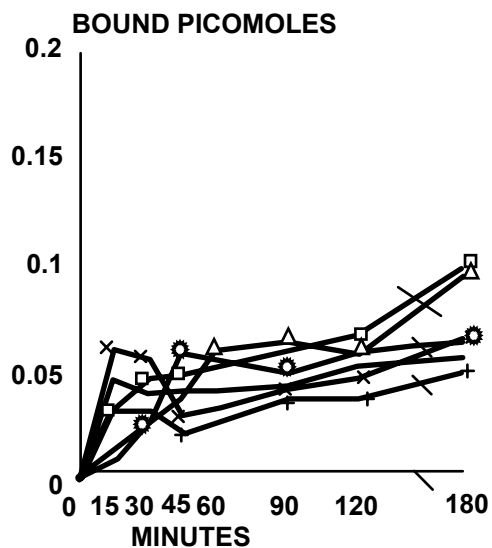
HIGH BINDING ACTIVITY PEPTIDES



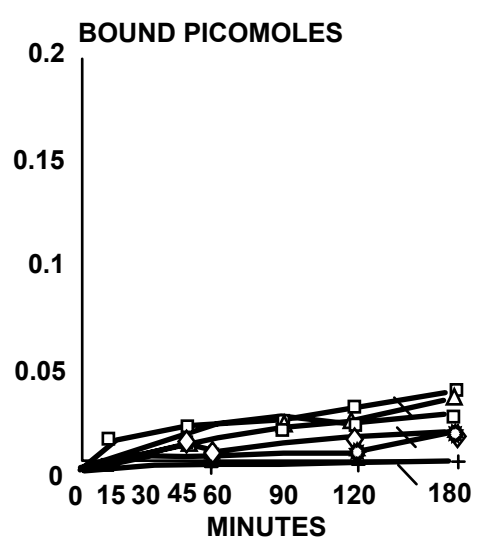
HIGH BINDING ACTIVITY PEPTIDES



MEDIUM BINDING ACTIVITY PEPTIDES



LOW BINDING ACTIVITY PEPTIDES





PRINCIPLES FOR VACCINE DEVELOPMENT

Receptor

Ligand

Host Cell

Parasite



INDUCE IMMUNITY



Blocking Receptor- Ligand



Vaccine

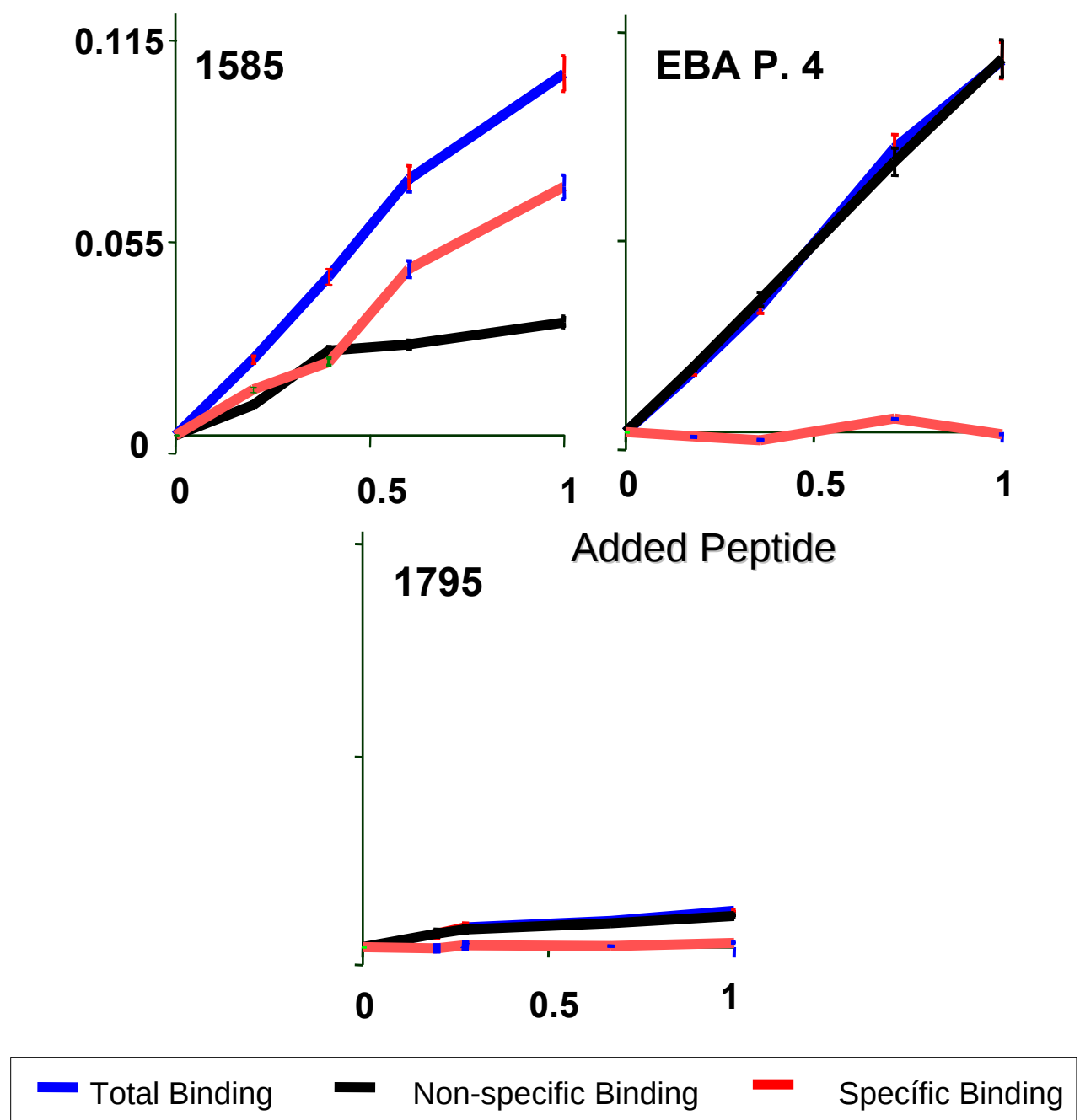


AMINO ACID SEQUENCE OF THE *EBA-175* SYNTHETIC PEPTIDES

Peptide number				Peptide number			
1755	20	KARNEYDIKENEKFLDVYKE	39	1791	740	TMCPEVKDVPISIIIRNNEQT	759
1756	40	KFNELDKKKYGNVQKTDKKI	59	1792	760	SQEAVPEENTEIAHRTETPS	779
1757	60	FTFIENKLDILNNSKFNKRW	79	1793	780	ISEGPKGNEQKERDDDSLK	799
1758	80	KSYGTPDNIDKNMSLINKHN	99	1794	800	ISVSPENSREPETAADKDTSNL	819
1759	100	NEEMFNNNYQSFLSTSSLIK	119	1795	820	LKLKGDVDISMPKAVIGSSP	839
1760	120	QNKYVPINAVRVSRILSFLD	139	1796	840	NDNINVTEQGDNISGVNSKP	859
1761	140	SRINNGRNTSSNNEVLSNCR	159	1797	860	LSDDVRPDKKELEDQNSDES	879
1762	160	EKRKGMKWDCCKKNDRSNYV	179	1798	880	EETVNHISKSPSINNGDDS	899
1763	180	CIPDRRIQLCIVNLSIIKTY	199	1799	900	SGSATVSESSSNTGLSID	919
1764	200	TKETMKDHFIEASKKESQLL	219	1800	920	DDRNGDTFVRTQDTANTEDV	939
1765	220	LKKNDNKYNSKFCNDLKNSF	239	1801	940	IRKENADKDEDEKGADEERH	959
1766	240	LDYGHLAMGNDMDFGGYSTK	259	1802	960	STSESLSSPEEKMLTDNEGG	979
1767	260	AENKIQEVFKGAHGEISEHK	279	1803	980	NSLNHEEVKEHTSNSDNVQQ	999
1768	280	IKNFRKEWWNEFREKLWEAM	299	1804	1000	SGGIVNMNVEKELKDTLENP	1019
1769	300	LSEHKNNINCKNIPQEELQ	319	1805	1020	SSSLDEGKAHEELSEPNLSS	1039
1770	320	ITQWIKEWHGFEFLERDNRS	339	1806	1040	DQDMSNTPGPLDNTSEETTE	1059
1771	340	KLPKSKCKNNTLYEACEKEC	359	1807	1060	RISNNEYKVNREDERTLT	1079
1772	360	IDPCMKYRDWIIRSKFEWHT	379	1808	1080	EYEDIVLKSHMNRESDDGEL	1099
1773	380	LSKEYETQKVPKENAENYLI	399	1809	1100	YDENSIDLSTVNDESEDAEAK	1119
1774	400	KISENKNDKAVSLLLNNCDA	419	1810	1120	MKGNDTSEMSHNSSQHIESD	1139
1775	420	EYSKYCDCKHTTTLVKSVLN	439	1811	1140	QQKNDMKTVDGLGTHVQNE	1159
1776	440	GNDNTIKEKREHIDLDDFSK	459	1812	1160	ISVPVTGEIDEKLRESKESK	1179
1777	460	FGCDKNSVDNTKVWECKNP	479	1813	1180	IHKAEERLSHTDIHKINPE	1199
1778	480	YILSTKDVCVPPRRQELCLG	499	1814	1200	DRNSNTLHLKDIRNEENERH	1219
1779	500	NIDRIYDKNLLMIKEHILAI	519	1815	1220	LTNQININISQERDLQKHGFH	1239
1780	520	AIYESRILKRKYKNKDDKEV	539	1816	1240	TMNNLHGDGVSESRQINHS	1259
1781	540	CKIINKTFADIRDIIGGTDY	559	1817	1260	HGNRQDRGGNSGNVLNMRN	1279
1782	560	WNDLSNRKLVGKINTNSKYV	579	1818	1280	NNNFNNIPSRYNLYDKKLDL	1299
1783	580	HRNKKNDKLFREWWKVIKK	599	1819	1300	DLYENRNDSTTKELIKKLAE	1319
1784	600	DVWNVISWVFKDKTVCKEDD	619	1820	1320	INKCENEISVKYCDHMIHEE	1339
1785	620	IENIPQFFRWFWSEWGDDYCQ	639	1821	1340	IPLKTCTKEKTRNLCCAVSD	1359
1786	640	DKTKMIETLKVECKEPCED	659	1822	1360	YCMSYFTYDSEYYNCTKRE	1379
1787	660	DNCKSKCNSYKEWISKKKEE	679	1823	1380	FDDPSYTCFRKEAFSSMIFK	1399
1788	680	YNKQAKQYQEYQKGNMY	699	1824	1400	FLITNKIYYFYTYKTAKVT	1419
1789	700	SEFKSIKPEVYLKKYSEKCS	719	1825	1420	IKKINFLIFFFFSF	1435
1790	720	NLNFEDEFKEELHSDYKNKC	739				

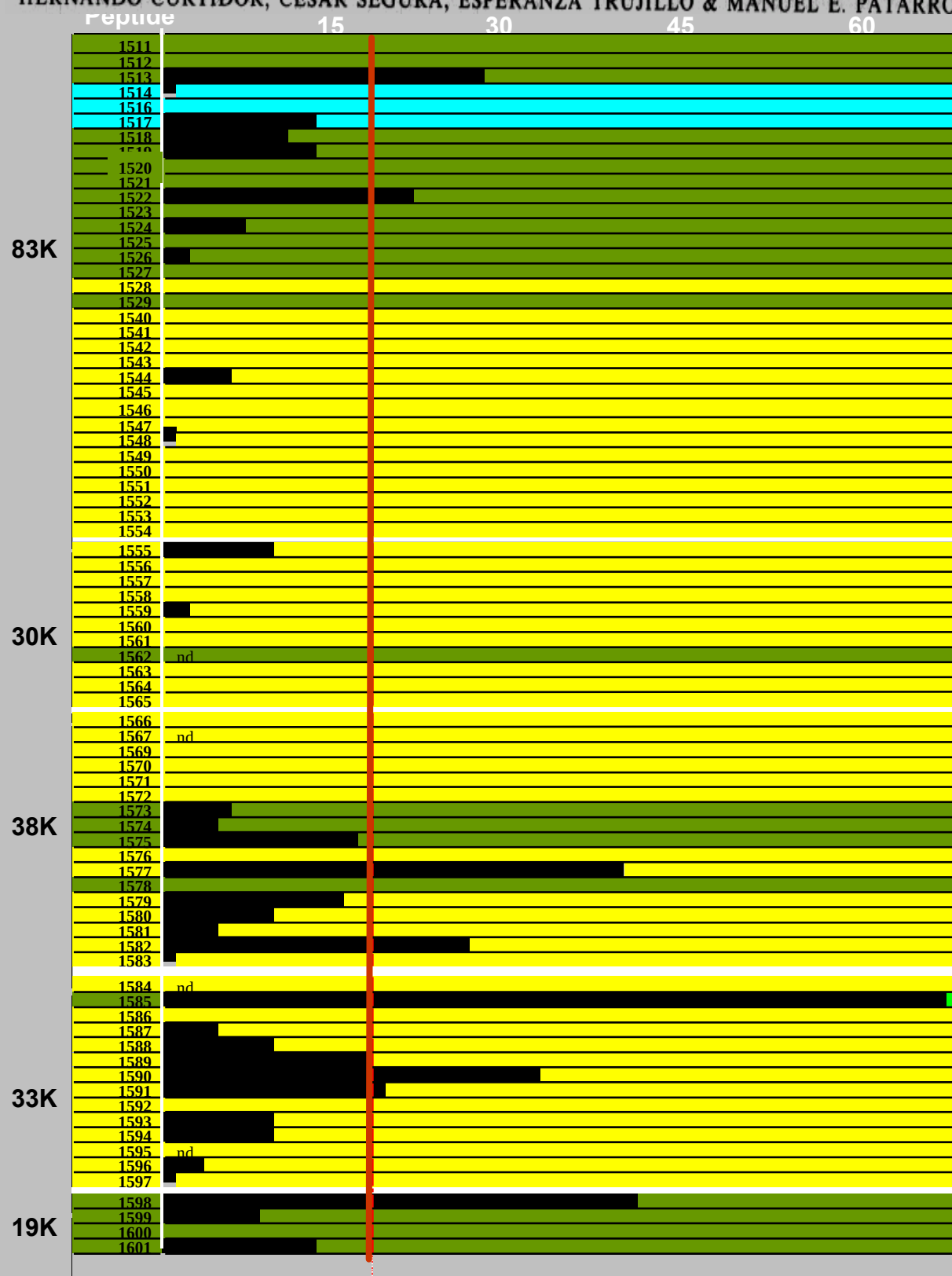
PEPTIDE RBC BINDING BEHAVIOUR

Peptide Binding activity is the curve's specific slope



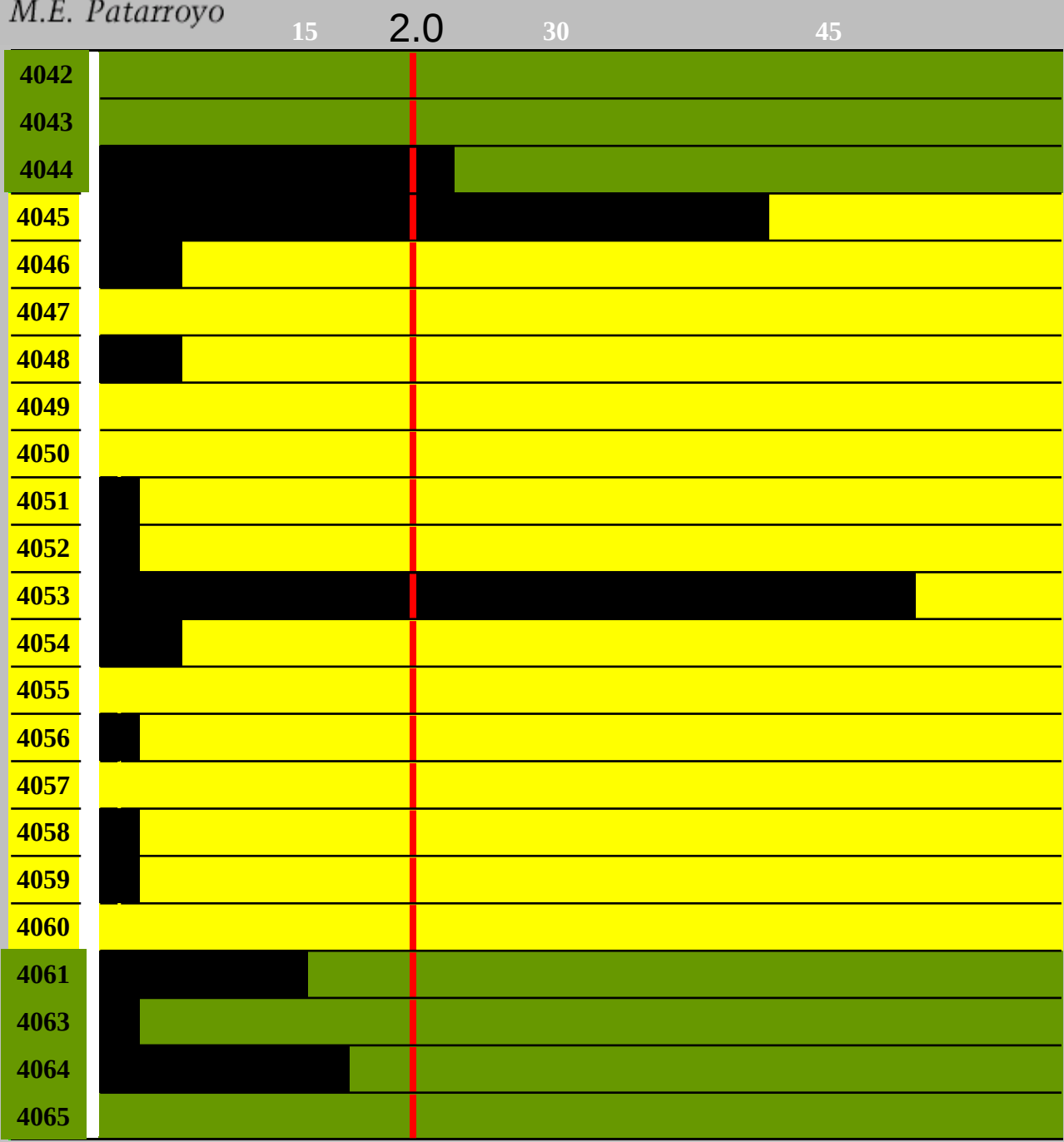
Identification of *Plasmodium falciparum* MSP-1 peptides able to bind to human red blood cells

MAURICIO URQUIZA, LUIS E. RODRIGUEZ, JORGE E. SUAREZ, FANNY GUZMÁN, MARISOL OCAMPO, HERNANDO CURTIDOR, CESAR SEGURA, ESPERANZA TRUJILLO & MANUEL E. PATARROYO



M. Ocampo
M. Urquiza
F. Guzmán
L.E. Rodríguez
J. Suárez
H. Curtidor
J. Rosas
M. Díaz
M.E. Patarroyo

Two MSA 2 peptides that bind
to human red blood cells are
relevant to *Plasmodium*
falciparum merozoite
invasion





Plasmodium falciparum normocyte binding protein (PfNBP-1) peptides bind specifically to human erythrocytes

John Jairo Valbuena, Ricardo Vera, Javier García, Alvaro Puentes, Hernando Curtidor, Marisol Ocampo, Mauricio Urquiza, Zuly Rivera, Fanny Guzmán, Elizabeth Torres, Manuel Elkin Patarroyo*

Normocyte Binding Protein PfNBP-1				Binding Activity	
Peptide		Sequence		1 %	2%
26327	1	LINIETEMKHKQKQLINKMNY	20		
26328	21	DIEKDNITDQYMHDVQQNIFY	40		
26329	41	EPITLKMNEYNTLLNDNHNN	60		
26330	61	NINNEHQFNHLNSLHTKIFSY	80		
26331	81	HNYNKEQQQEYITNIMQRID	100		
26332	101	VFINDLDTYQYEEYFYEWNQ	120		
26333	121	EYKQIDKNKINQHINNINNN	140		
26334	141	LIHVKKQFEHTLENIKNNENY	160		
26335	161	IFDNIQLKKKDIDDIININY	180		
26336	181	NTKETYLKELNKKKMLQNKK	200		
26337	201	KVDEKSEINNHHHTLQHDNQNY	220		
26338	221	VEQKNKIKDHNLTCPNNNSY	240		
26339	241	SEESHQNEQMKEQNKNILEKY	260		
26340	261	QTRNIKPHHVHNHNHNHNHNY	280		
26341	281	HNQNNQKQDSTKLQEEDISTY	300		
26342	301	HKLHNTIHEQQSKDNHQGNRY	320		
26343	321	EKKQKNGNHERMYFASGIVV	340		
26344	341	SILFLSSLGFVINSKNNKQEY	360		
26345	361	YDKEQEKQQQNDNFVCDNNKM	380		
26346	381	DDKSTQKYGRNQEEVMEISF	400		
26347	386	QKYGRNQEEVMEISFDNDYI	405		

AMA-1

Urquiza M., *et al.*, 2001, *Vaccine*, 19: 508



FIDIC

Peptide

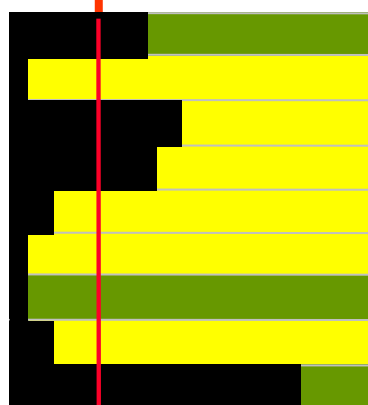
Sequence

2.0

4307	14	EFTYMINFGRGQNYWEHPYQ	33
4308	34	KSDVYHPINEHREHPKEYQY	53
4309	54	PLHQEHTYQQEDSGEDENTL	73
4310	74	QHAYPIDHEGAEPAPQEQL	93
4311	94	FSSIEIVERSNYMGNPWTEY	113
4312	114	MAKYDIEEVHSGSIRVDLGE	133



4313	134	DAEVAGTQYRLPSGKCPVFG	153
4314	154	KGIIENSNTTFLTPVATGNY	173
4315	174	QYLKDGGFAPPTTEPLMSPM	193
4316	194	TLDEMRHFYKDNKYVKNLDE	113
4317	214	LTLC SRHAGNMI PDNDKNSNY	233
4318	234	YKPAVYDDKDKKCHILYIA	253
4319	254	AQENNGPRYCNKDESKRNSM	273
4320	274	FCFRPAKDISFQNYTYLSKN	293
4321	294	VVDNWEKVCPRKNLQNAKFGY	313



Peptide

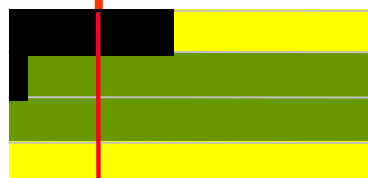
Kd (nM)

4313	120 ± 12
4315	120 ± 10
4316	150 ± 14
4322	700 ± 21
4325	100 ± 09
4328	140 ± 11

4322	314	LWDGNCEDI PHVNEFSAIDY	333
4323	334	LFECNKL VFELSASDQPKQY	353
4324	354	EQHLTDYEKIKEGFKNKNAS	373
4325	374	MIKSAFLPTGAFKADRYKSH	393
4326	394	GKGYNWGNYNTETQKCEIFN	413
4327	414	VKPTCLINNSSYIATTALSH	433



4328	434	PIEVEHNFPCLYKNEIMKE	453
4329	454	IERESKR I KLNDNDDEGNKK	473
4330	474	IIAPRIFISDDKDSLKCPCDY	493
4331	494	PEIVSNSTCNFFVCKCVERIFY	513



4332	514	AEVTSNNEVVVKEEYKDEYA	533
4333	534	DIP EHKPTYDKMKII IASSA	553
4334	554	AVAVLAT I LMVYLYKRKGNA	573
4335	574	EKYDKMDEPQHYGKSNSRND	593
4336	594	EMLDPEASFWGEEKRASHTTY	613
4337	603	WGEEKRASHTTPVLMEKPYY	622



Plasmodium falciparum EBA-175 kDa protein peptides which bind to human red blood cells

L. E. RODRIGUEZ*, M. URQUIZA, M. OCAMPO, J. SUAREZ, H. CURTIDOR, F. GUZMAN, L. E. VARGAS, M. TRIVIÑOS, M. ROSAS and M. E. PATARROYO*



ELSEVIER

Vaccine 18 (2000) 1289–1293

Vaccine

Amino terminal peptides of the ring infected erythrocyte surface antigen of *Plasmodium falciparum* bind specifically to erythrocytes

Ricardo Vera Bravo, Viviana Marín, Javier García, Mauricio Urquiza, Elizabeth Torres, Mary Trujillo, Jaiver Rosas, Manuel Elkin Patarroyo*



ELSEVIER

Parasitology International 49 (2000) 105–117



Serine repeat antigen peptides which bind specifically to red blood cells

A. Puentes*, Javier Garcia, Ricardo Vera, Q. Ramses Lopez, Mauricio Urquiza, Magnolia Vanegas, Luz Mary Salazar, Manuel Elkin Patarroyo

L.E. Rodriguez
M. Ocampo
R. Vera
A. Puentes
R. Lopez
J. Garcia
H. Curtidor
J. Valbuena
J. Suarez
J. Rosas
Z. Rivera
M. Urquiza
M.E. Patarroyo

Plasmodium falciparum
EBA-140 kDa protein
peptides that bind to human
red blood cells

J. Peptide Res., 2003, 61, 175–184.

Plasmodium falciparum: red blood cell binding studies using peptides derived from rhoptry-associated protein 2 (RAP2)

Ramsés López ^{a,b}, John Valbuena ^{a,b}, Hernando Curtidor ^{a,b}, Alvaro Puentes ^{a,b}, Luis E. Rodríguez ^{a,b}, Javier García ^{a,b}, Jorge Suárez ^{a,b}, Ricardo Vera ^{a,b}, Marisol Ocampo ^{a,b}, Mary Trujillo ^{a,b}, Luis E. Ramírez ^{a,b}, Manuel E. Patarroyo ^{a,b,*}

Plasmodium falciparum: red blood cell binding studies of peptides derived from histidine-rich KAHRP-I, HRP-II and HRP-III proteins

Ramsés López ^{*}, Mauricio Urquiza, Hernando Curtidor, Jorge Eduardo Caminos, Hernando Mora, Alvaro Puentes, Manuel Elkin Patarroyo

Plasmodium falciparum acid basic repeat antigen (ABRA) peptides: erythrocyte binding and biological activity

H. Curtidor, M. Urquiza, J.E. Suarez, L.E. Rodriguez, M. Ocampo, A. Puentes, J.E. Garcia, R. Vera, R. López, L.E. Ramirez, M. Pinzon, M.E. Patarroyo ^{*}

Identification of specific Hep G2 cell binding regions in *Plasmodium falciparum* sporozoite–threonine–asparagine-rich protein (STARP)

Ramsés López, Javier Garcia, Alvaro Puentes, Hernando Curtidor, Marisol Ocampo, Ricardo Vera, Luis Eduardo Rodriguez, Jorge Suarez, Mauricio Urquiza, Ana Liliana Rodriguez, Claudia Alexandra Reyes, Carmen Giovana Granados, Manuel E. Patarroyo ^{*}

R. López
H. Curtidor
M. Urquiza
J. Garcia
A. Puentes
J. Suarez
M. Ocampo
R. Vera
L.E. Rodriguez
F. Castillo
G. Cifuentes
M.E. Patarroyo

Plasmodium falciparum: binding studies of peptide derived from the sporozoite surface protein 2 to Hep G2 cells

J. Peptide Res., 2001, 58, 285–292.



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Vaccine xxx (2003) xxx–xxx

Vaccine

www.elsevier.com/locate/vaccine

Sporozoite Liver Stage Antigen peptides bind specifically to human hepatocytes

Alvaro Puentes, Javier García, Ricardo Vera, Ramsés López, Jorge Suarez, Luis Rodriguez, Hernando Curtidor, Marisol Ocampo, Diana Tovar, Martha Forero, Adriana Bermudez, Jimena Cortes, Mauricio Urquiza, Manuel E. Patarroyo*



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Parasitology International xx (2003) xxx–xxx



www.elsevier.com/locate/parint

Identification of *Plasmodium falciparum* reticulocyte binding protein RBP-2 homologue a and b (PfRBP-2-Ha and -Hb) sequences that specifically bind to erythrocytes

Marisol Ocampo¹, Ricardo Vera¹, Luis Eduardo Rodríguez, Hernando Curtidor, Jorge Suárez, Javier García, Alvaro Puentes, Ramsés López, John Valbuena, Diana Tovar, Claudia Reyes, Sandra Vega, Manuel Elkin Patarroyo*



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Vaccine xxx (2003) xxx–xxx

Vaccine

www.elsevier.com/locate/vaccine

Specific erythrocyte binding capacity and biological activity of *Plasmodium falciparum*-derived rhoptry-associated protein 1 peptides

Hernando Curtidor, Marisol Ocampo, Diana Tovar, Ramses López, Javier García, Jhon Valbuena, Ricardo Vera, Jorge Suárez, Luis E. Rodríguez, Álvaro Puentes, Fanny Guzmán, Elizabeth Torres, Manuel E. Patarroyo*



Vaccine 19 (2001) 508–513

Vaccine

www.elsevier.com/locate/vaccine

Plasmodium falciparum AMA-1 erythrocyte binding peptides implicate AMA-1 as erythrocyte binding protein

Mauricio Urquiza, Jorge E. Suarez, Constanza Cardenas, Ramses Lopez, Alvaro Puentes, Francisco Chavez, Julio Cesar Calvo, Manuel Elkin Patarroyo*

Instituto de Inmunología, Hospital San Juan de Dios, Universidad Nacional de Colombia, AA 44709 Bogotá, Colombia



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Biochimie des protéines

BIOCHIMIE

Plasmodium falciparum: red blood cell binding studies using peptides derived from rhoptry-associated protein 2 (RAP2)

Ramsés López ^{a,b}, John Valbuena ^{a,b}, Hernando Curtidor ^{a,b}, Alvaro Puentes ^{a,b}, Luis E. Rodríguez ^{a,b}, Javier García ^{a,b}, Jorge Suárez ^{a,b}, Ricardo Vera ^{a,b}, Marisol Ocampo ^{a,b}, Mary Trujillo ^{a,b}, Luis E. Ramirez ^{a,b}, Manuel E. Patarroyo ^{a,b,*}



Vaccine 19 (2001) 4487–4495

Vaccine

Plasmodium falciparum circumsporozoite (CS) protein peptides specifically bind to HepG2 cells

Jorge E. Suarez ^{*}, Mauricio Urquiza, Alvaro Puentes, Javier E. Garcia, Hernando Curtidor, Marisol Ocampo, Ramses Lopez, Luis E. Rodriguez, Ricardo Vera, Marcia Cubillos, Maria H. Torres, Manuel E. Patarroyo



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Peptides xxx (2003) xxx–xxx

PEPTIDES

www.elsevier.com/locate/peptides

P. falciparum: merozoite surface protein-8 peptides bind specifically to human erythrocytes

Alvaro Puentes, Javier García, Marisol Ocampo, Luis Rodríguez, Ricardo Vera, Hernando Curtidor, Ramsés López, Jorge Suarez, John Valbuena, Magnolia Vanegas, Fanny Guzman, Diana Tovar, Manuel E. Patarroyo^{*}



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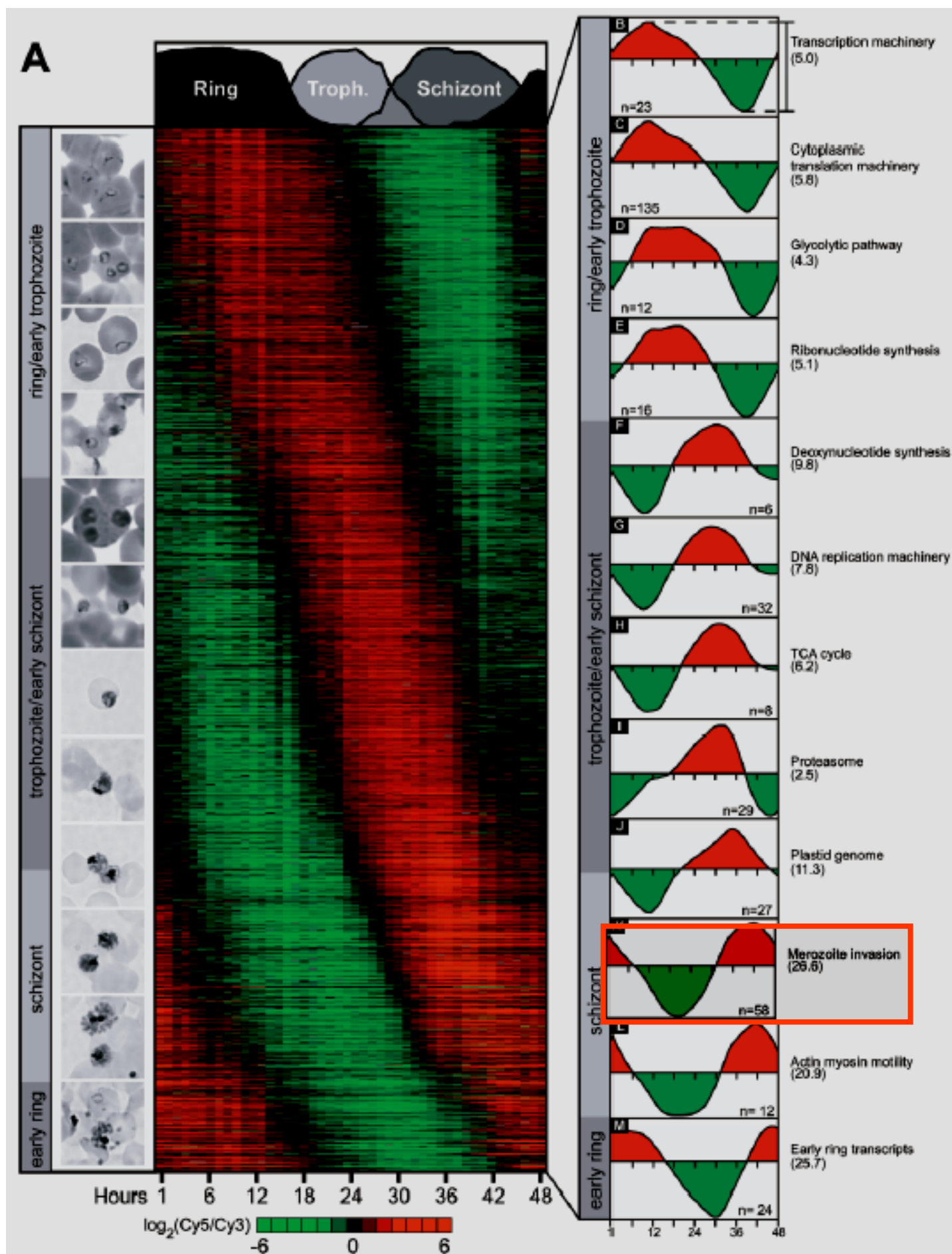
Peptides 24 (2003) 647–657

PEPTIDES

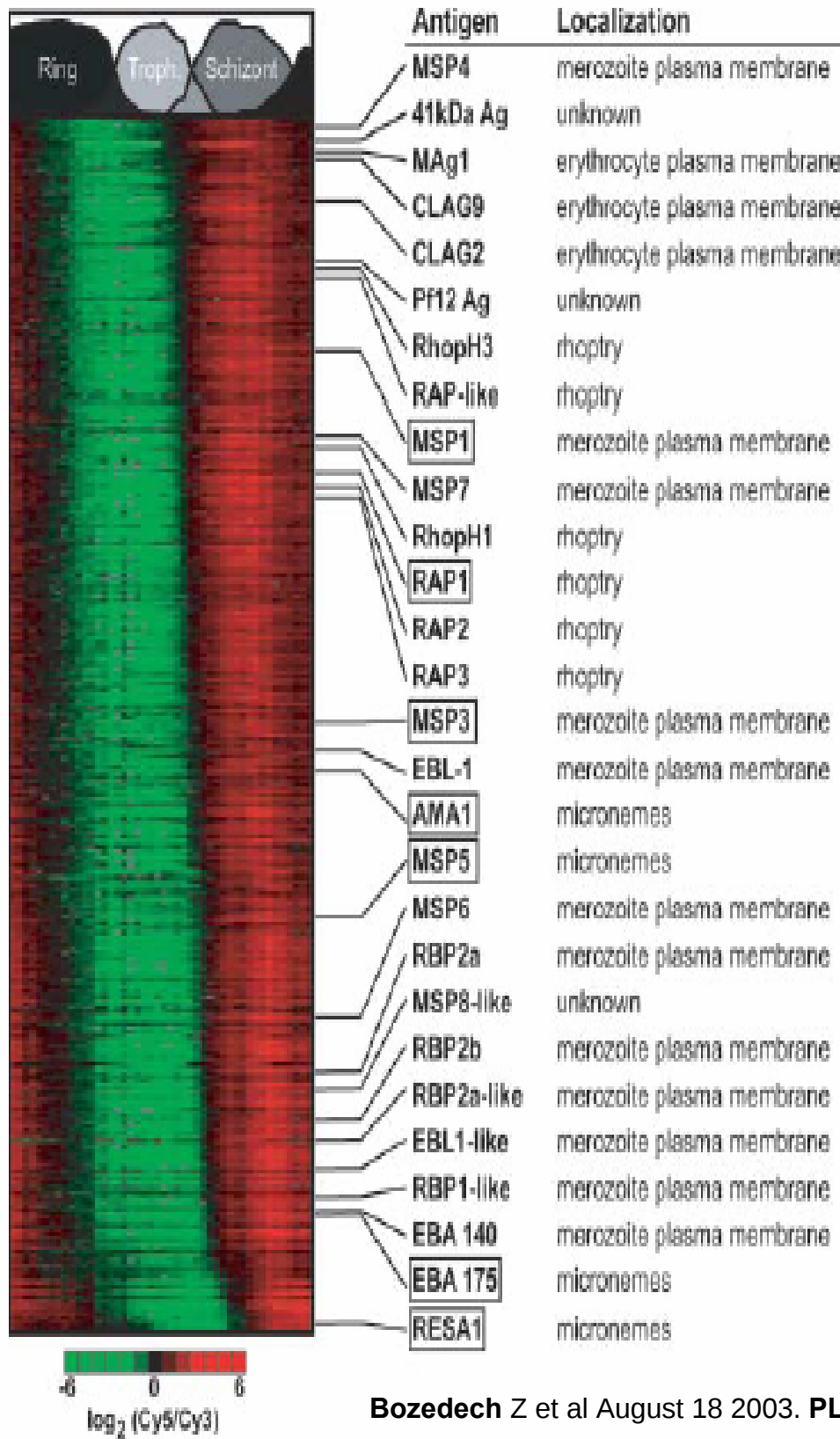
www.elsevier.com/locate/peptides

Peptides of the liver stage antigen-1 (LSA-1) of *Plasmodium falciparum* bind to human hepatocytes

Javier E. García, Alvaro Puentes, Ramsés López, Ricardo Vera, Jorge Suárez, Luis Rodríguez, Hernando Curtidor, Marisol Ocampo, Diana Tovar, Martha Forero, Adriana Bermudez, Jimena Cortés, Mauricio Urquiza, Manuel E. Patarroyo^{*}



PHASEOGRAM OF PUTATIVE VACCINE TARGETS



Bozdech Z et al August 18 2003. **PLOS Biology** 1, 001

High RBC binding activity peptides



FIDIC

PROTEIN	C	V	Reference
MSP1	3	6	<u>Urquiza M. et. al . 1996. <i>Parasite Immunology</i> . 18: 515-526</u>
RESA	2	–	<u>Vera R. et al . 2000. <i>Vaccine</i>. 18: 1289-1293</u>
SERA	6	1	<u>Puentes A. et. al . 2000. <i>Parasitology International</i>. 49: 105-117</u>
MSA2	1	2	<u>Ocampo M. et. al . <i>J. Peptide Research</i>. 55: 216-233</u>
EBA175	6	–	<u>Rodríguez L.E et. al. 2000. <i>Parasitology</i>. 120: 225-235</u>
GBP130	1	–	<u>Suárez J. et. al . 2000. <i>Mem. Inst. Oswaldo Cruz</i> . 95: 495-501</u>
HRP I, II, II	3	1	<u>López R. et a l. 2000. <i>Acta Tropica</i>. 75: 349-359</u>
ABRA	5	–	<u>Curtidor H. et. al . 2001. <i>Vaccine</i>. 19: 4496-4504</u>
AMA1	4	3	<u>Urquiza M. et. al . 2001. <i>Vaccine</i>. 19: 508-513</u>
EBA140	6	–	<u>Rodríguez L.E.. et. al . 2003. <i>J. Peptide Research</i>. 62: 175-184</u>
NBP1	2	–	<u>Valbuena J. et al . 2003. <i>Peptides</i> . 24: 1007-1014</u>
MSP8	5	–	<u>Puentes A. et. al . 2003. <i>Peptides</i> .24: 1015-1023</u>
MAEBL	9	–	<u>Ocampo M. et al . 2004. <i>Biochem. Biophys. Res. Comm</i>. 315: 319-329</u>
RAP2	4	–	<u>López R. et al . 2004. <i>Biochemie</i> . 86: 1-6</u>
RBP2 Ha/Hb	9	3	<u>Ocampo M. et. al . 2004. <i>Parasitol. Int</i>. 53: 77-88</u>
RAP1	4	–	<u>Curtidor H. et. al . 2004. <i>Vaccine</i> . 22: 1054-1062</u>
EBA160	5	–	<u>Valbuena J. et al . 2004. <i>Biochem. Biophys. Res. Comm</i> . 321: 835-844</u>
CLAG3	5	–	<u>Ocampo M. et. al . 2005. <i>Proteins Science</i> . 14: 504-513</u>
EBL1	5	–	<u>Curtidor H. et. al . 2005. <i>Protein Science</i> . 14: 464-473</u>
JESEBL	5	–	<u>Vera R. et al . 2004. <i>Biochemie. In press</i></u>
STEVOR	–	3	<u>García J. et al . 2004. <i>Peptides. In press</i></u>
MSP10	3	–	<u>Puentes A. et al . 2004. <i>Biochemie. In press</i></u>
MSP3	3	–	<u>Rodríguez L.E. et. al . 2004. <i>Submitted to Protein Science</i></u>
SORTILIN	6	–	<u>Vera R. et al . 2004. <i>Submitted to Biochemistry</i></u>
RESA-LIKE	3	–	<u>Rodríguez L.E. et. al . 2004. <i>Submitted to J. Int. Parasitol.</i></u>
TryThrA	4	–	<u>Curtidor H. et. al . 2004. <i>Submitted to Chembiochem</i></u>
HAP	2	–	<u>Valbuena J. et. al . 2004. <i>Submitted to Biol. Chem.</i></u>
MSP6	2	1	<u>López R. et al . 2004. <i>Submitted to Peptides</i></u>

C : CONSERVED

V: VARIABLE

Synthetic vaccines - Emerging principles

- 1. Plasmodia (falciparum & vivax) use CONSERVED as well as VARIABLE sequences to bind to host cells.**

ANTIGENICITY STUDIES

HYPERIMMUNE MONKEYS



Aotus Monkeys develop resistance to reinfection after being infected several times with $5 \text{ to } 500 \times 10^6$ *P. falciparum* FVO infected erythrocytes and treated with chloroquine

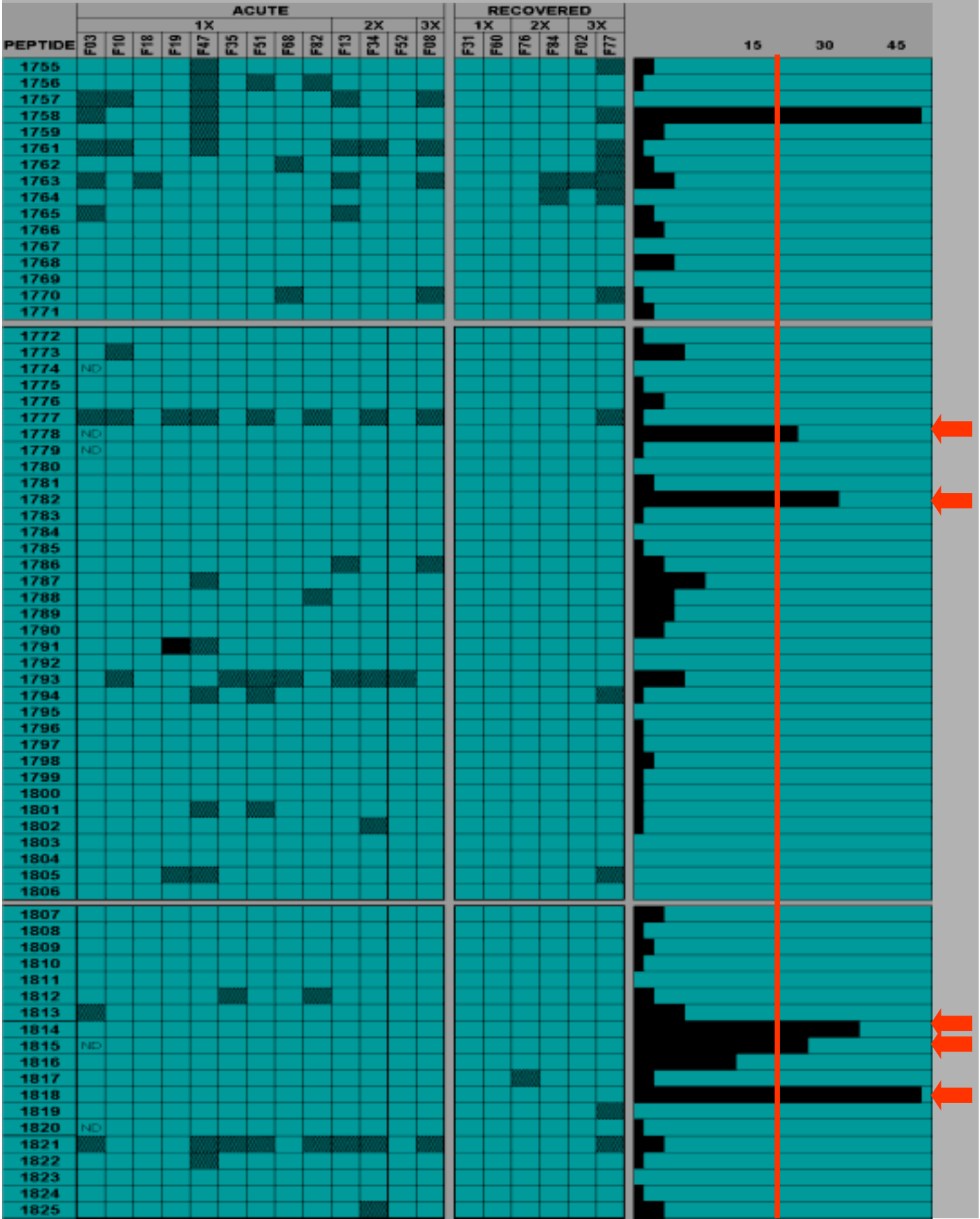
AOTUS ABS AGAINST MSP-2 PEPTIDES Vs. BINDING ACTIVITY

[illegible]

HUMAN ABS AGAINST MSP-2 PEPTIDES Vs. BINDING ACTIVITY

83K

HUMAN Abs AGAINST EBA-175 PEPTIDES vs. BINDING



Synthetic vaccines - Emerging principles

1. Plasmodia (falciparum & vivax) use **CONSERVED** as well as **VARIABLE** sequences to bind to host cells.
2. **CONSERVED** binding sequences are **NOT** antigenic, but **VARIABLE** sequences are

IMMUNOGENICITY AND PROTECTION STUDIES

PROTOCOL FOR IMMUNIZATION OF AOTUS MONKEYS



Monkeys were immunized with 125 μ gr of peptide in FA on days 1, 20, and 40, bled on day 60 for immunological studies, and challenge on day 65.

AOTUS SERA REACTIVITY AGAINST ORIGINAL EBA-175 IMMUNIZING PEPTIDES

EBA
175 Kd

GROUP H	1678	1692	1713	1730	1737
1758	0	0	0	200	0
1779	0	0	0	0	0
1783	0	100	0	100	0

EBA 175 Kd
VARIABLE

GROUP I	1679	1693	1714	1731	1738
1814	100	200	12800	6400	6400
1815	12800	25600	51200	12800	1600

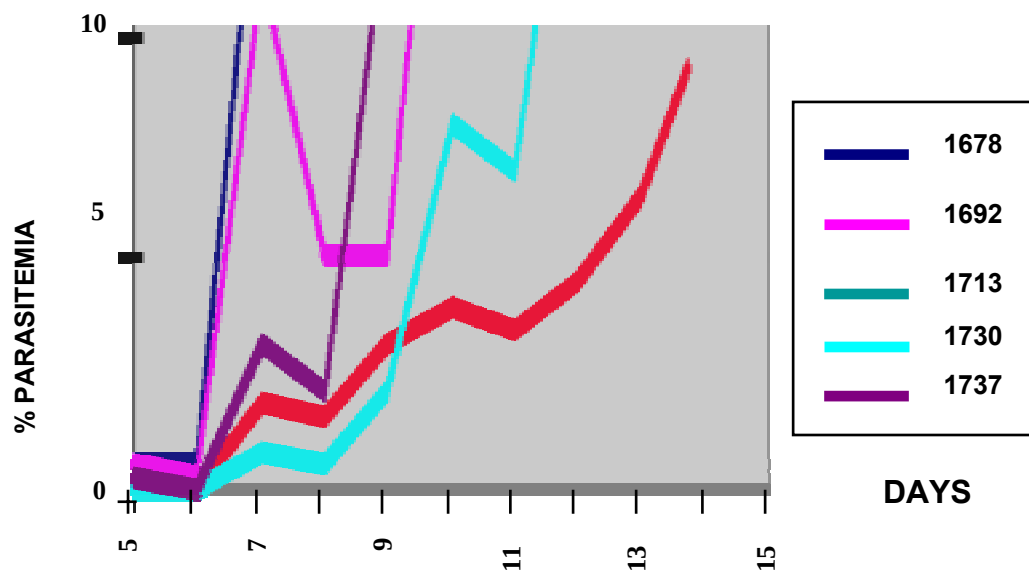
EBA 175 Kd
C&V

GROUP J	1681	1694	1715	1732	1739
1758	0	0	0	400	100
1779	0	0	0	0	0
1783	0	0	0	800	0
1814	200	100	100	100	0
1815	25600	204800	25600	3200	6400

AOTUS MONKEYS IMMUNIZED WITH EBA UNMODIFIED PEPTIDES

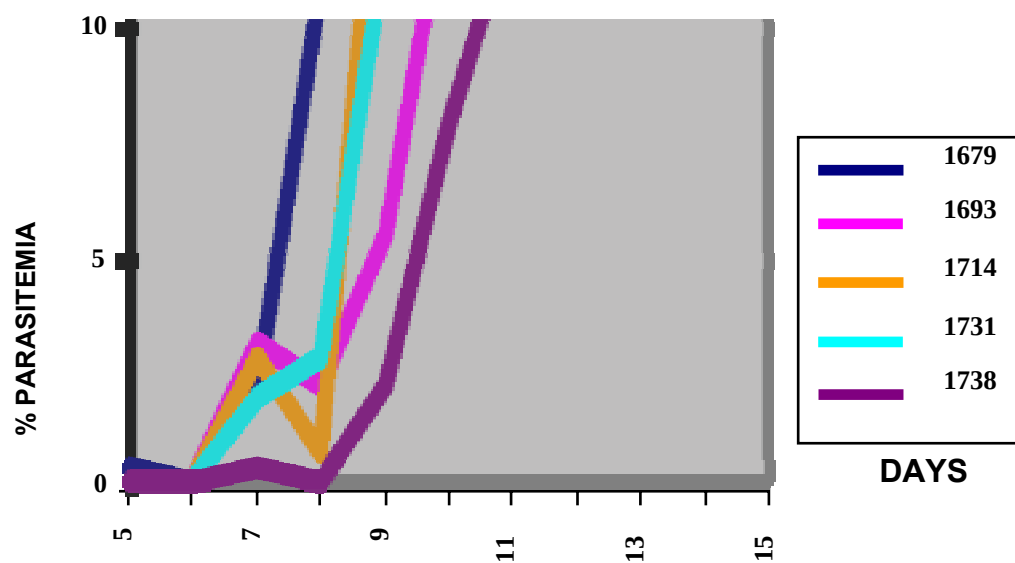
GROUP H: IMMUNIZED WITH **CONSERVED** HBAP

1758, 1779, 1783.



GRUPO I: IMMUNIZED WITH **VARIABLE** HBAP

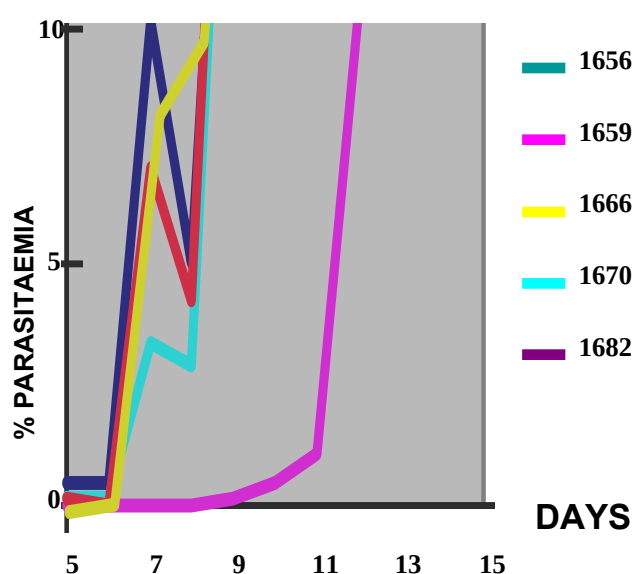
1814, 1815.



AOTUS MONKEYS IMMUNIZED WITH MSP-1 UNMODIFIED PEPTIDE MIXTURES

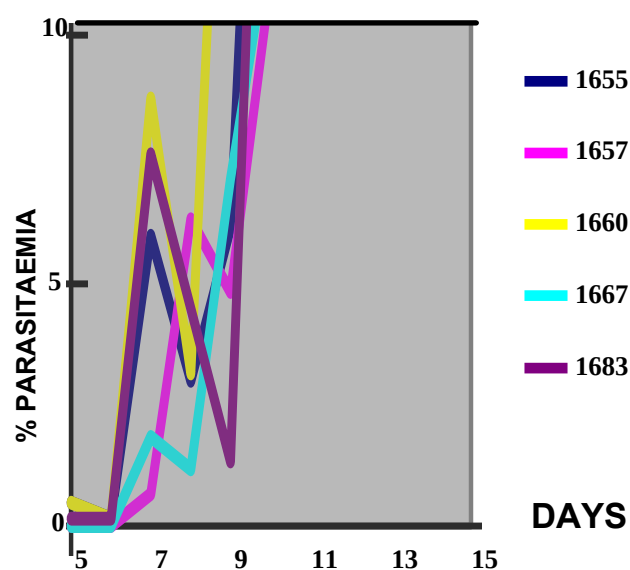
GROUP A: IMMUNIZED WITH PEPTIDES FROM

83 Kd :
1513, 1517, 1522



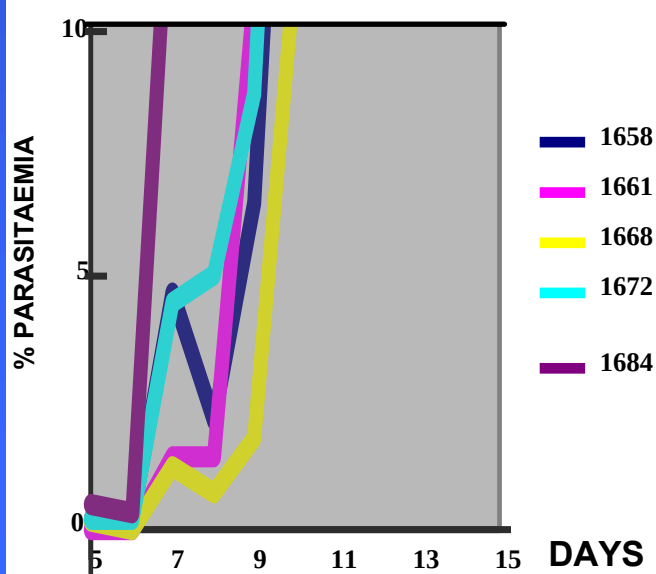
GROUP B: IMMUNIZED WITH PEPTIDES FROM

38 Kd :
1577, 1582.



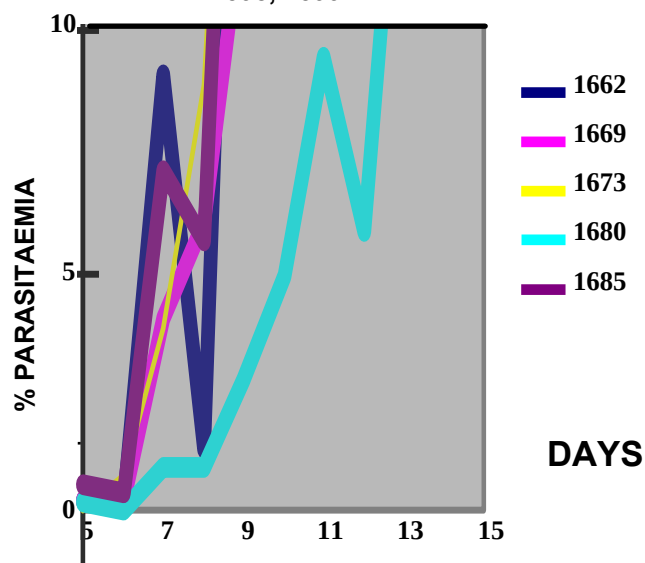
GROUP C: IMMUNIZED WITH PEPTIDES

FROM **33 Kd:**
1585, 1590, 4017.



GROUP D: IMMUNIZED WITH PEPTIDES

FROM **19 Kd:**
1598, 1599.



MAPS USED IN THE MONKEY TRIALS

MAP 5A 1512 (H E S Y Q E L V K K L E A L E D A V L T) ₄ - P₃C

MAP 5B 1513 (G Y S L F Q K E K M V L N E) ₄ - P₃C

MAP 6A 1522 (Q I P F N L K I R A N E L D V L K K L V) ₄ - P₃C

MAP 6B 1523 (F G Y R K P L D N I K D N V G K M E D Y) ₄ - P₃C

MAP 8B 1584 (L G Q V V T G E A V T P S V I D N I L S K I E N E Y) ₄ - P₃C

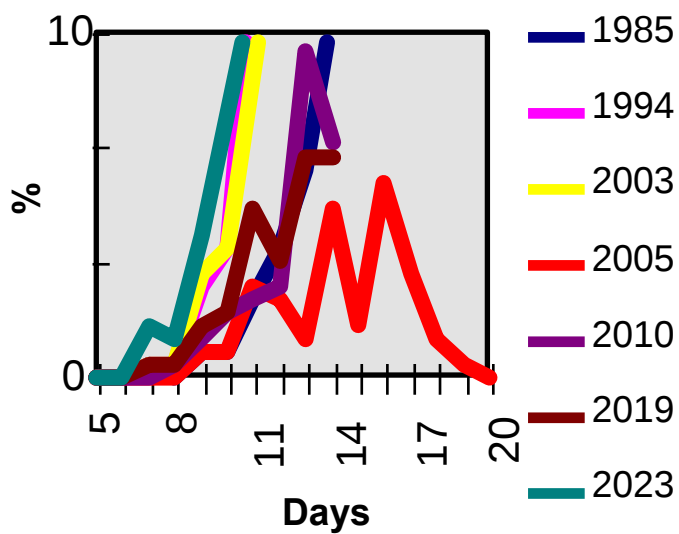
MAP 9B 1585 (D V I Y L K P L A G V Y R S L K K Q I E) ₄ - P₃C

MAP 11A 1596 (I A D L S T D Y N H N N L L T K F L S T G) ₄ - P₃C

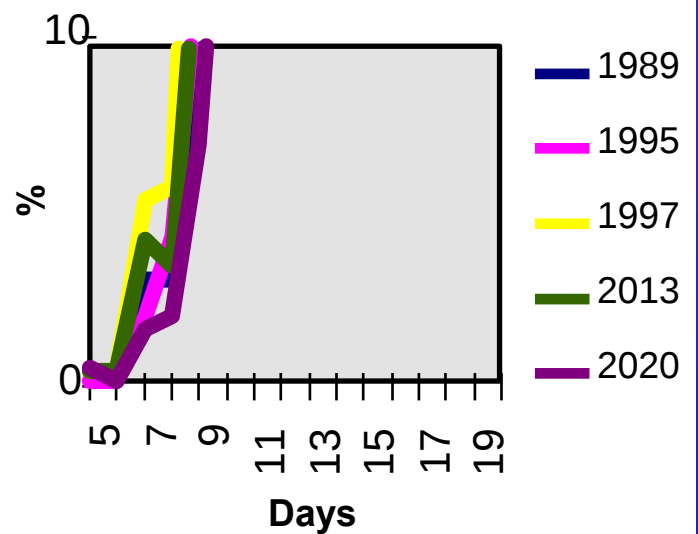
MAP 11B 1597x (M V F E N L A K T V L S N L L D N L Q G M L N I S Q H Q) ₄ - P₃C

AOTUS MONKEYS IMMUNIZED WITH MAP MIXTURES

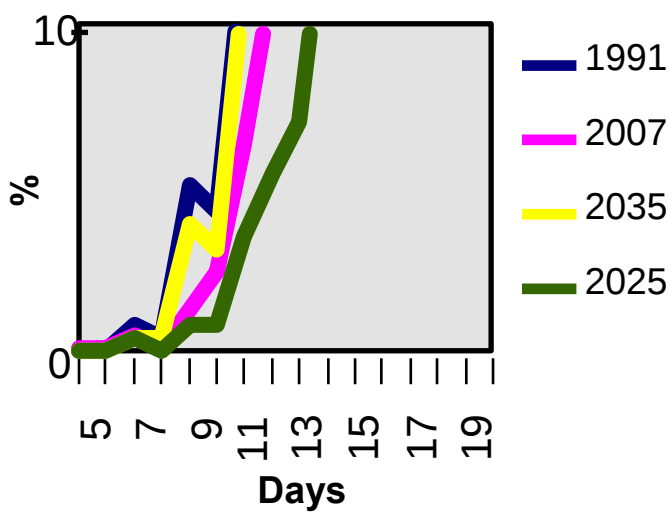
MAPS 3X



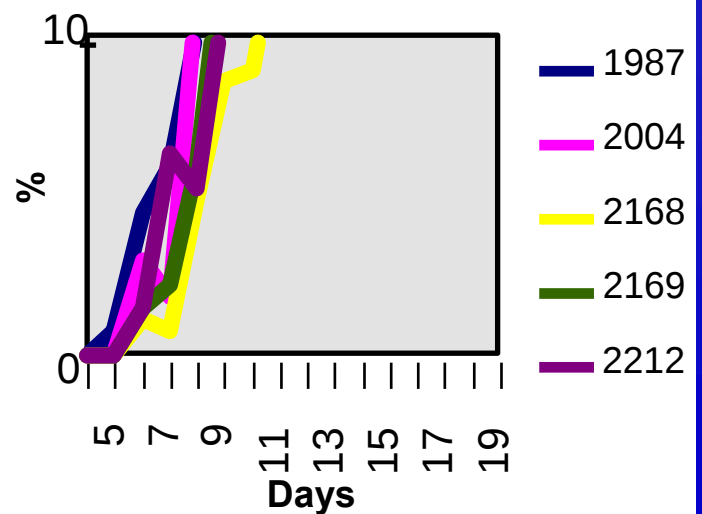
MAPS 5X



CONTROLS



CONTROLS



Aminoacid Sequence of the MSP-1, EBA-175, GBP-130, and MSP-2 Expanded Peptides

Patarroyo M. E. et al. Submitted to *J. Structural Biology*

1512	HESYQELVKKLEALED AVL	1513	TGYSLFQKEK MVLNE	A
1522	QIPFNLKIRANELDVLKKLV	1523	FGYRKPLDNIKDNV GKMEDY	B
1576	LKTLSEESIQTEDNYASLEN	1577	FKVLSKLEGK LKDNLNLEKK	C
1582	AESNTITTSQNVDDEVDDVILGQ	1584	VVTGEAVTPSVIDNILSKIENEY	D
1584	LDQVVTGEAISVTMDNILSGFENEY	1585	DVIY LKPLAGVYRSLKKQIE	E
1596	IADLSTDYNHNNLLTKFLSTGM	1597	VFENLAKTVLSNLLDNLQGMLNISQHQ	G
1757	FTFIENKLDILNNSKF NKRW	1758	KSYGTPDNIDKNNSLINKHN	H
1779	LGNIDRIYDKNLLMIKEHILAIAI	1780	YESRILKRKYKNKDDKEV	I
1782	YWNDLSNRKLVGKINTNSK	1783	YVHRNKKNDK LFRDEWWKVIK D	J
1814	DRNSNTLHLKDIRNEEMERH	1815	LTNQINISQERDLQKHGFH	K
2208	IFHKILTNTDPNDEVERRNA	2209	ADPEYRKHLEV FHKILTNTD	L
2219	EGQIMREYASDPEYRKHLEI	2220	FYKILTNTDPND DVERRNAD	M

AOTUS SERA REACTIVITY AGAINST EBA-175 EXPANDED PEPTIDES

Patarroyo M. E. et al. Submitted to *J. Structural Biology*

GROUP E

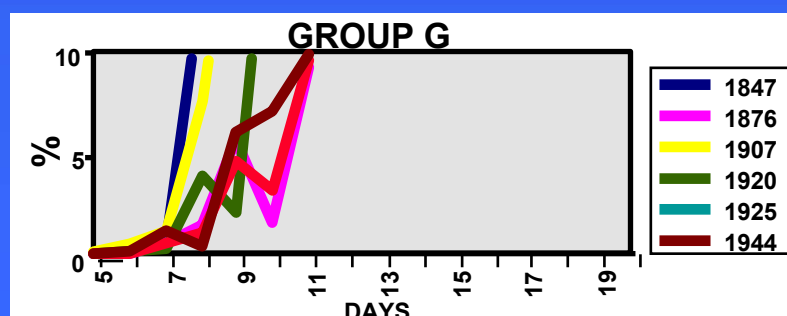
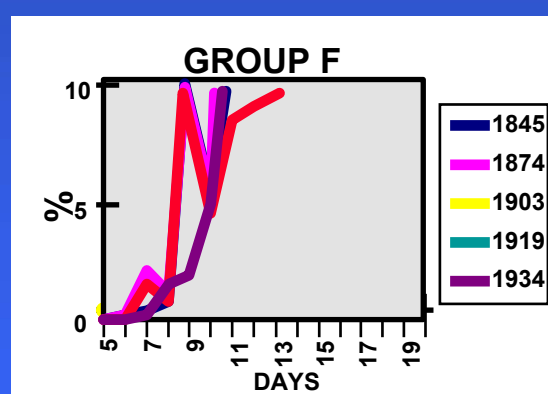
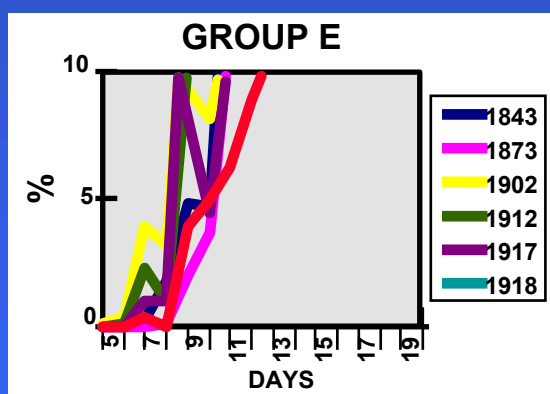
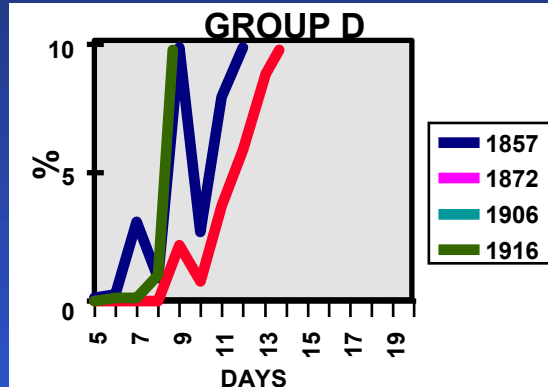
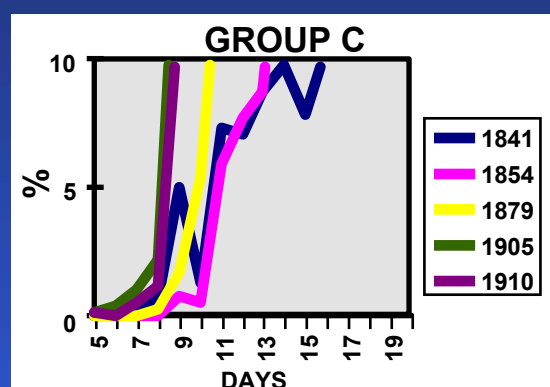
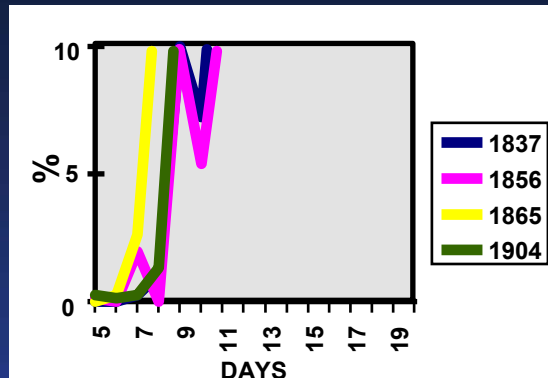
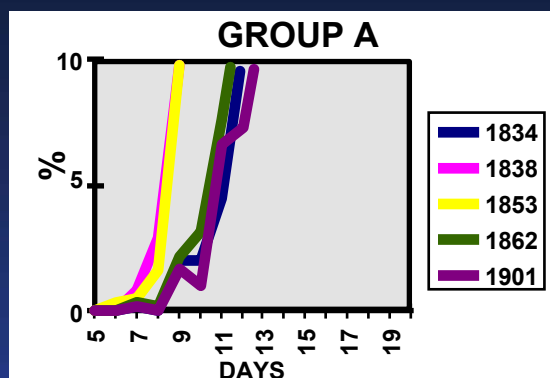
	1843	1873	1902	1912	1917	1918
1779	0	0	0	0	0	0
1780	800	0	6400	0	100	1600
1782	12800	800	3200	12800	12800	400
1783	0	0	100	100	200	100

GROUP F

	1845	1874	1903	1919	1934
1757	25600	3200	0	12800	0
1758	0	100	0	100	0
1814	400	100	100	200	800
1815	51200	6400	6400	25600	6400

AOTUS MONKEYS IMMUNIZED WITH MSP-1 EXPANDED PEPTIDES

Patarroyo M. E. et al. Submitted to *J. Structural Biology*



IMMUNOGENICITY OF THE BINDING PEPTIDES IN RABBITS

CONSERVED				DIMORPHIC		
	PEPTIDE Nr.	RABBIT Nr.	Ab TITER	PEPTIDE Nr.	RABBIT Nr.	Ab TITER
MSA-1	1513	182	1600	1589	205	12800
	1522	190	0	1525	207	204800
	1577	180	800	1590	193	1600
	1585	189	0	1561	228	102400
	1598	181	0	1565	218	6400
MSP-2	4045	195	800	4053	192	12800
EBA-175	1758	200	800	1814	199	25600
	1779	185	800	1815	184	204800
	1783	186	0	1818	222	25600
				1756	209	102400
				1782	210	12800
GBP-130	2220	172	1600	2202	204	12800
				2204	216	12800

IMMUNOGENICITY OF THE CONSERVED BINDING PEPTIDES



For the production of Mo Abs groups of 4 Balb/c mice were immunized i.p. 5x with 15 μ gr of the individual MSP1 (1513, 1522, 1585, 5501) EBA - 175 (1758, 1779, 1783, 1818) GBP - 130 (2220) or MSP₂ (4044) conserved peptides. After the 3rd immunization NOT EVEN A SINGLE MOUSE, showed antibodies against the corresponding immunizing peptide, as assessed by ELISA western blots or IFA. The same was also true for mice of the AKR and DBA strains.

Synthetic vaccines - Emerging principles

1. Plasmodia (falciparum & vivax) use CONSERVED as well as VARIABLE sequences to bind to host cells.
2. CONSERVED binding sequences are NOT antigenic, but VARIABLE sequences are
3. **CONSERVED binding sequences are NOT immunogenic in ANY of the tested species when administered as monomers, polymerized, associated with T helper epitopes or presented as MAPs. While VARIABLE sequences can be immunogenic they are strain-specific**

AN EMPIRICAL APPROACH

Changing some residues.

- **To convert putative alpha helices into beta II or beta VII turns.**
- **To modify putative aromatic composition.**

STRUCTURALLY REPLACED ANALOGUES OF THE 1513 PEPTIDE

PEPTIDE	GROUP	SEQUENCE
1513	1 2 3	TGYSLFQKEKMVLNEGTS GTA
4863		TG M SL P QKEKMVLNE
4865		TG H SLFQKEKMVLNE
4887		TGYSL H QKEKMVLNE

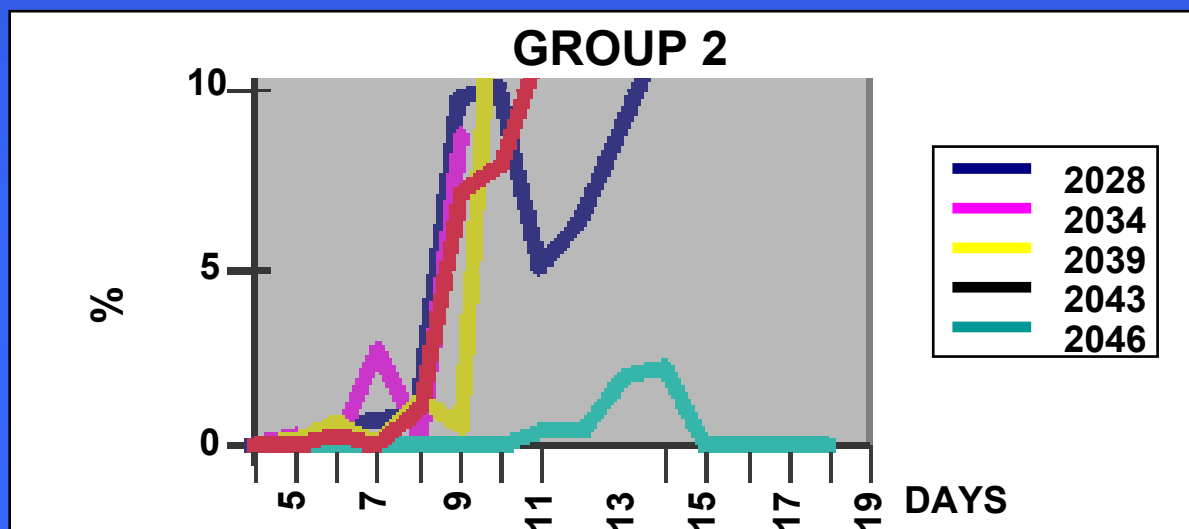
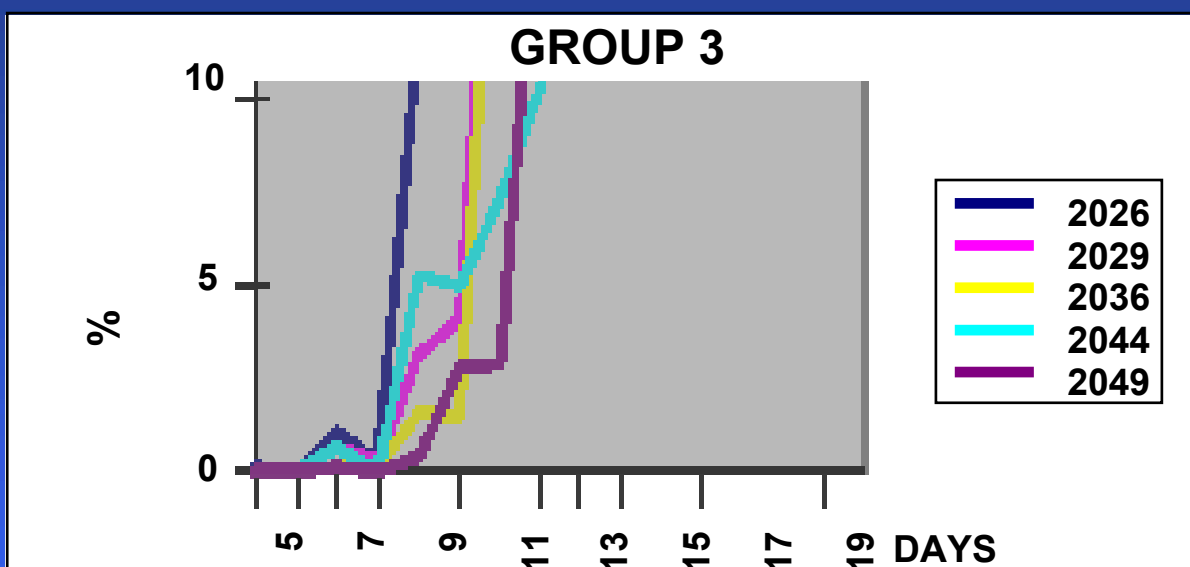
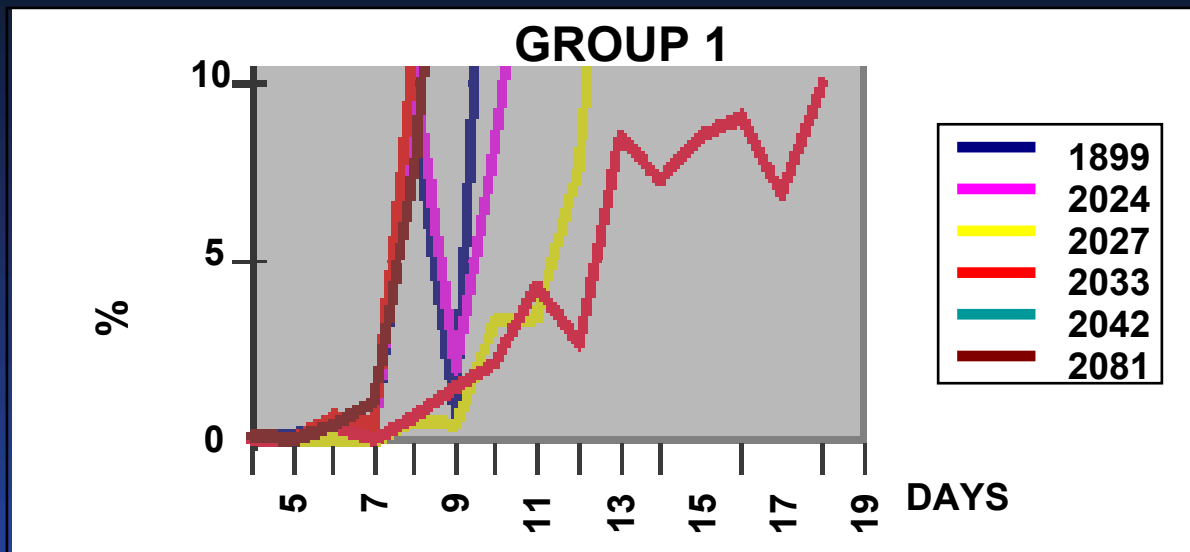
AOTUS SERA REACTIVITY AGAINST THE STRUCTURALLY REPLACED IMMUNIZING PEPTIDES

GROUP 1	2024	2027	2033	2042	2081
1513	0	0	0	0	0
4863	0	0	0	0	0
4865	0	0	0	0	100
4867	0	0	0	0	0

GROUP 2	2028	2034	2039	2043	2046
1513	800	0	0	1600	400
4863	0	0	0	25600	0
4865	800	100	0	3200	200
4867	0	0	0	6400	6400

GROUP 3	2026	2029	2036	2044	2049
1513	0	200	0	1600	100
4863	0	100	0	6400	0
4865	200	100	0	6400	0
4867	200	200	200	12800	0

AOTUS MONKEYS IMMUNIZED WITH MSP-1 STRUCTURALLY MODIFIED PEPTIDES



STRUCTURALLY REPLACED ANALOGUES OF THE 1598_M PEPTIDE

PEPTIDE	GROUP	SEQUENCE
4847	14	FPNTIISKLIIEGKFQDMLNISQHQ
4871	4	M PN P IISKLIIEGKFQDMLNISQHQ
4873	5	FPNTIISKLIIEGKFQDM P N P SQHQ
4879	6	FPNTIISKLIIEGK P QDM P N P SQHQ

AOTUS SERA REACTIVITY AGAINST THE STRUCTURALLY REPLACED IMMUNIZING PEPTIDES

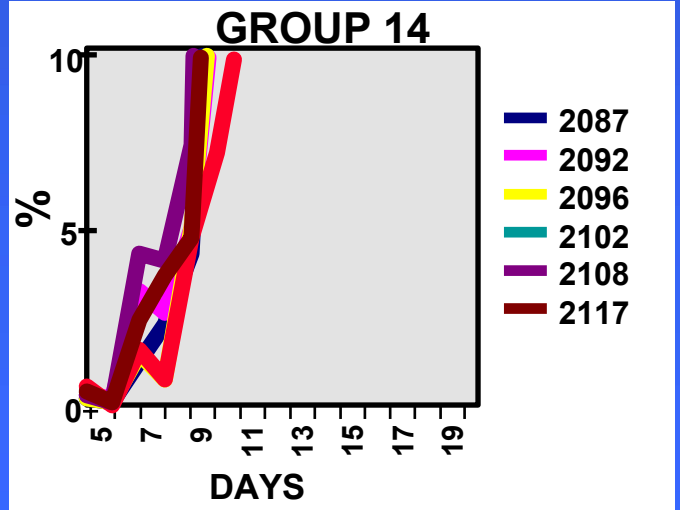
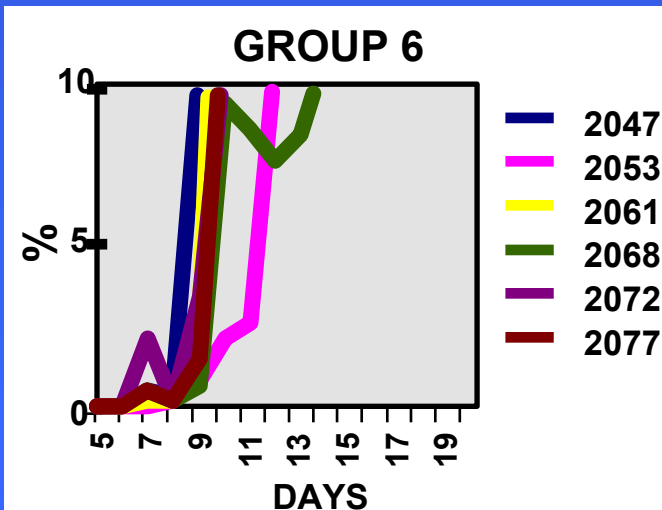
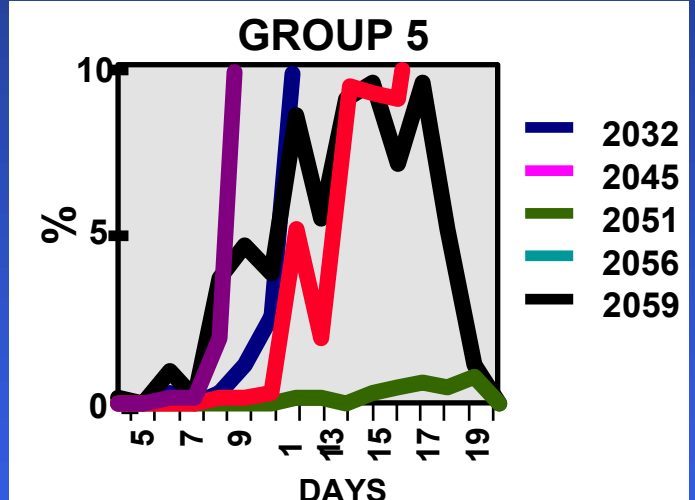
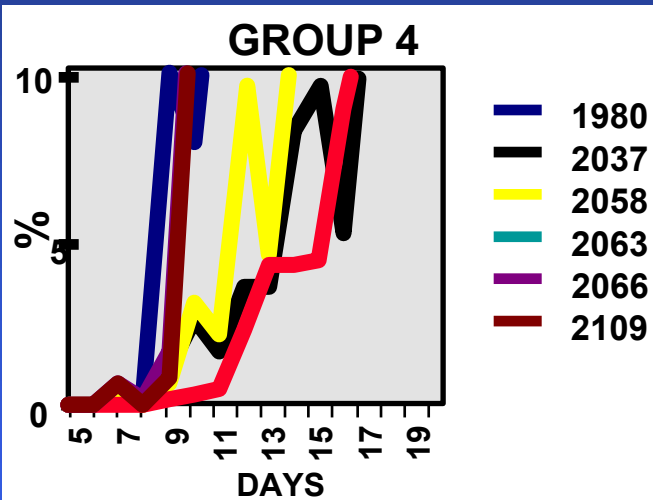
GROUP 4	1980	2037	2058	2063	2066	2109
4869	100	200	200	12800	51200	200
4871	1600	102400	800	102400	102400	25600
4873	100	ND	0	ND	ND	200
4879	3200	12800	6400	51200	6400	3200

GROUP 5	2032	2045	2051	2056	2059	2047
4869	100	3200	1600	51200	3200	200
4871	200	800	200	1600	200	25600
4873	200	ND	ND	ND	ND	ND
4879	51200	51200	12800	12800	25600	12800

GROUP 6	2053	2061	2068	2072	2077
4869	51200	102400	12800	25600	102400
4871	25600	51200	3200	6400	12800
4873	ND	ND	ND	ND	ND
4879	51200	25600	800	102400	204800

GROUP 14	2087	2092	2096	2102	2108	2117
4869	0	0	0	0	0	0
4871	0	0	0	0	0	0
4873	0	0	0	0	0	0
4879	0	0	0	0	0	0

AOTUS MONKEYS IMMUNIZED WITH MSP-1 STRUCTURALLY MODIFIED PEPTIDES



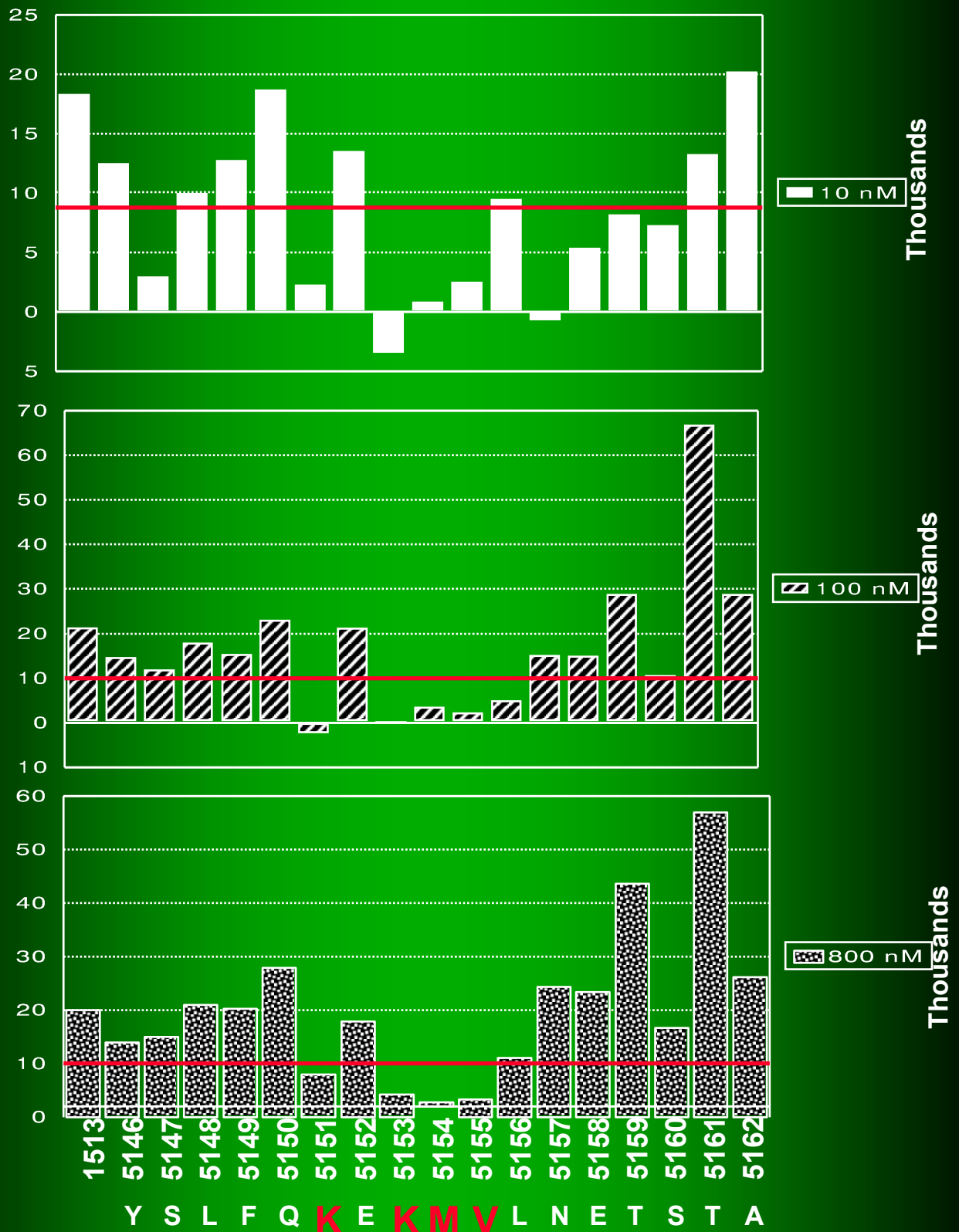
ANALOGUES OF PEPTIDE 1513 FOR BINDING ASSAYS

Urquiza M. et al. 1996. *Parasite Immunology* 18:515-526

1513	G	Y	S	L	F	Q	K	E	K	M	V	L	N	E	G	T	S	G	T	A	Y
5146	G	G	S	L	F	Q	K	E	K	M	V	L	N	E	G	T	S	G	T	A	Y
5147	G	Y	G	L	F	Q	K	E	K	M	V	L	N	E	G	T	S	G	T	A	Y
5148	G	Y	S	G	F	Q	K	E	K	M	V	L	N	E	G	T	S	G	T	A	Y
5149	G	Y	S	L	G	Q	K	E	K	M	V	L	N	E	G	T	S	G	T	A	Y
5150	G	Y	S	L	F	G	K	E	K	M	V	L	N	E	G	T	S	G	T	A	Y
5151	G	Y	S	L	F	Q	G	E	K	M	V	L	N	E	G	T	S	G	T	A	Y
5152	G	Y	S	L	F	Q	K	G	K	M	V	L	N	E	G	T	S	G	T	A	Y
5153	G	Y	S	L	F	Q	K	E	G	M	V	L	N	E	G	T	S	G	T	A	Y
5154	G	Y	S	L	F	Q	K	E	K	G	V	L	N	E	G	T	S	G	T	A	Y
5155	G	Y	S	L	F	Q	K	E	K	M	G	L	N	E	G	T	S	G	T	A	Y
5156	G	Y	S	L	F	Q	K	E	K	M	V	G	N	E	G	T	S	G	T	A	Y
5157	G	Y	S	L	F	Q	K	E	K	M	V	L	G	E	G	T	S	G	T	A	Y
5158	G	Y	S	L	F	Q	K	E	K	M	V	L	N	G	G	T	S	G	T	A	Y
5159	G	Y	S	L	F	Q	K	E	K	M	V	L	N	E	G	G	S	G	T	A	Y
5160	G	Y	S	L	F	Q	K	E	K	M	V	L	N	E	G	T	G	G	T	A	Y
5161	G	Y	S	L	F	Q	K	E	K	M	V	L	N	E	G	T	S	G	G	A	Y
5162	G	Y	S	L	F	Q	K	E	K	M	V	L	N	E	G	T	S	G	T	G	Y

BINDING OF 1513 PEPTIDE ANALOGUES TO RBCs

Urquiza M. et al. 1996. *Parasite Immunology* 18:515-526



SPECIFICALLY REPLACED ANALOGUES OF THE 1585 PEPTIDE TO IMMUNIZE AOTUS MONKEYS

Espejo F. et al. 2001. *Angewandte Chemie Int. Ed.* 40:2631

PEPTIDE

SEQUENCE

1585 EVLY**LKP****L**AGVYRSLKKQLE

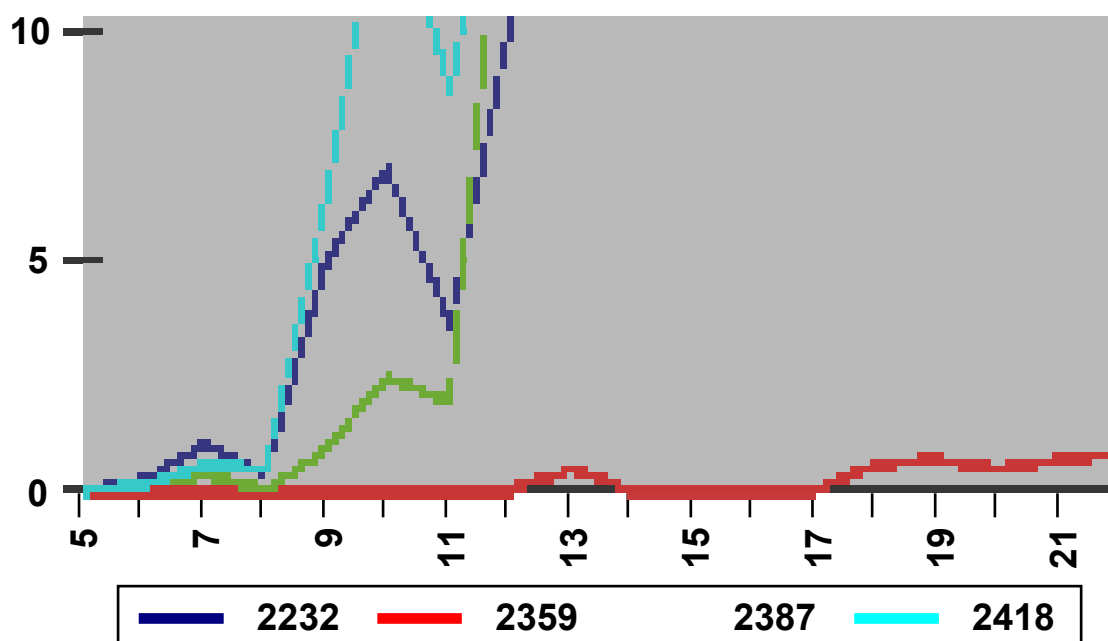
5187 EVLY**G**KPLAGVYRSLKKQLE

5190 EVLYLKP**G**AGVYRSLKKQLE

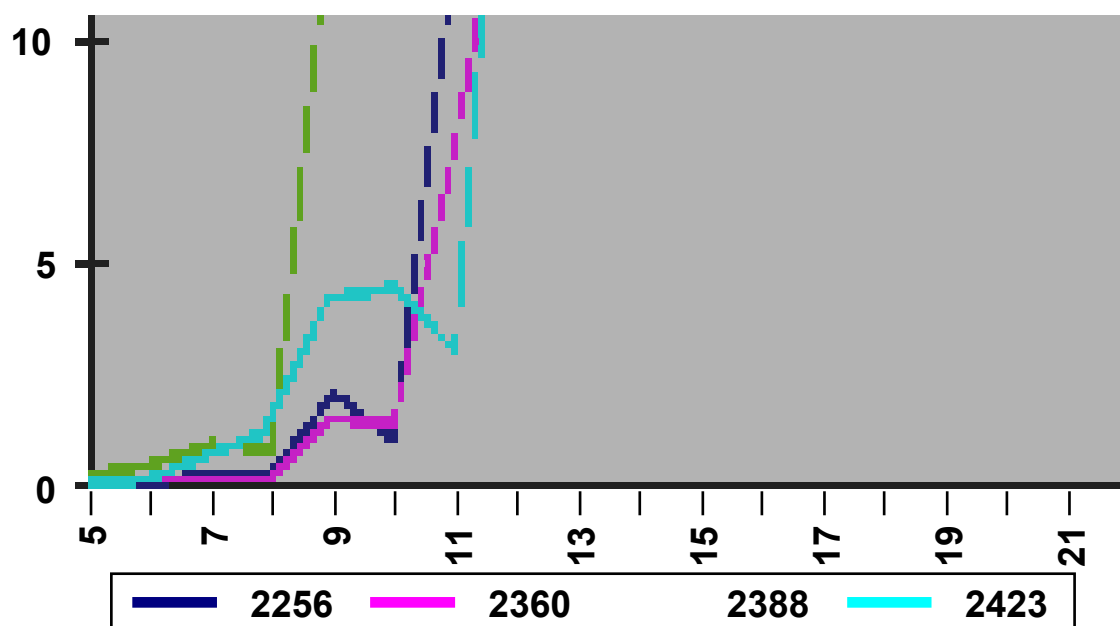
AOTUS MONKEYS IMMUNIZED WITH 1585 SPECIFICALLY MODIFIED PEPTIDES

Espejo F. et al. 2001. *Angewandte Chemie Int. Ed.* 40:2631

GROUP 7



GROUP 8



SPECIFICALLY REPLACED ANALOGUES OF THE 5501 PEPTIDE TO IMMUNIZE AOTUS MONKEYS

Torres M. H. et al. 2003. *European J. Biochemistry* 270:3946

PEPTIDE

SEQUENCE

5501 MLNISQHQCVKKQCPQNSY

5503 MGNISQHQCVKKQCPQNSY

5512 MLNISQHQCVGKQCPQNSY

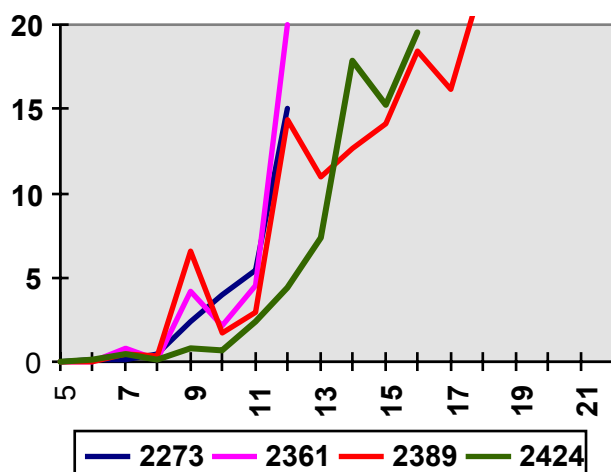
5513 MLNISQHQCVKGQCPQNSY

5516 MLNISQHQCVKKQCGQNSY

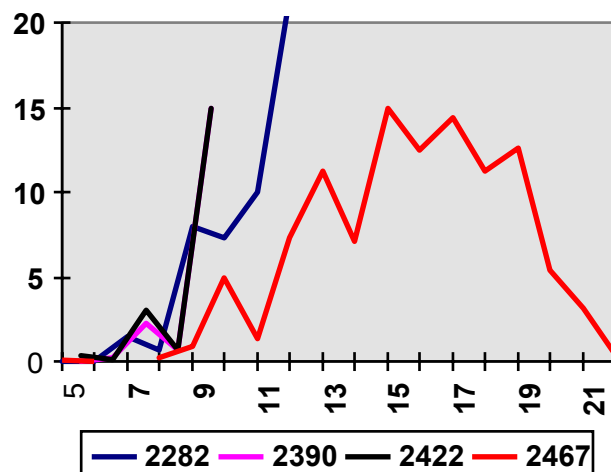
AOTUS MONKEYS IMMUNIZED WITH 5501 SPECIFICALLY MODIFIED PEPTIDES

Torres M. H. et al. 2003. *European J. Biochemistry* 270:3946

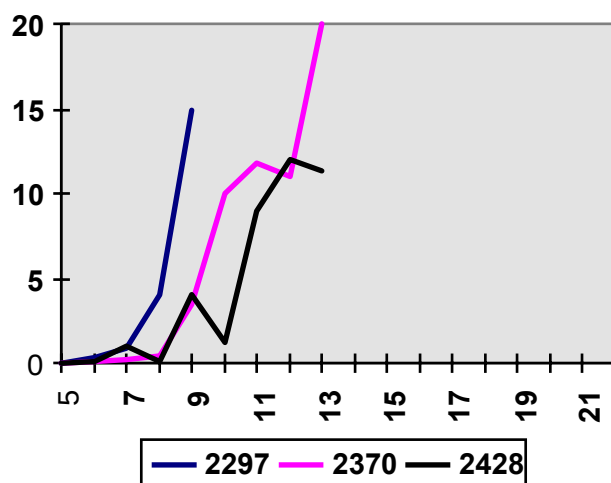
GROUP 9



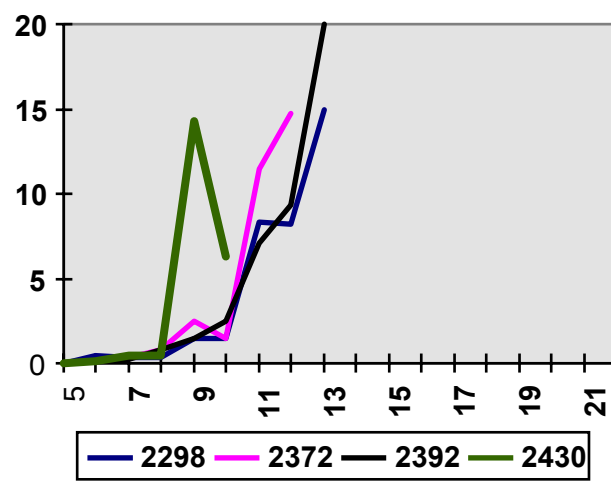
GROUP 10



GROUP 11



GROUP 12



Synthetic vaccines - Emerging principles

1. Plasmodia (falciparum & vivax) use CONSERVED as well as VARIABLE sequences to bind to host cells.
2. CONSERVED binding sequences are NOT antigenic, but VARIABLE sequences are
3. CONSERVED binding sequences are NOT immuno-genic in ANY of the tested species when administered as monomers, polymerized, associated with T helper epitopes or presented as MAPs. While VARIABLE sequences can be immunogenic they are strain-specific
4. To overcome this problem the critical contact residues in the binding interaction have to be identified and changed

COMMUNICATIONS

Structure, Immunogenicity, and Protectivity Relationship for the 1585 Malarial Peptide and Its Substitution Analogues

Fabiola Espejo,* Marcia Cubillos, Luz Mary Salazar, Fanny Guzman, Mauricio Urquiza, Marisol Ocampo, Yolanda Silva, Raul Rodriguez, Eduardo Liroy, and Manuel Elkin Patarroyo*

Polymerized peptide Nr.	Peptide Sequence	IFA TITERS ≥ 320		
		II 15	III 20	PROT
1585	E V L Y L K P L A G V Y R S L K K Q L E	0	0	0/5
5187	- - - - G K P A - - - - S - - - - -	ND.	1(320)	1/4
5188	- - - - L G P A - - - - S - - - - -	ND.	1(320)	1/4
5189	- - - - L K G A - - - - S - - - - -	ND.	1(1280)	1/4
6187	- - - - L K P G A - - - S - - - - -	0	0	0
6177	- - - - L K P A - - - - G - - - - -	0	0	0
13450	- - - - L L D A - - - - S - - - - -	2(2560)	1(2560)	2/4
22806	- - - - L L D A - - - - S - - - - -	3(1280)	1(1280)	2/9
10014	- - - - H V P A - - - - S - - - - -	2(640)	1(640)	2/4
22768	- - - - H L P A - - - - A - - - - -	0	1(5120)	1/10
11860	- - - - H M P G - - - - A - - - - -	1(1280)	1(2560)	2/6
22770	- - - - H L P G - - - - A - - - - -	1(1280)	1(1280)	2/9
13454	- - - - L M S A - - - - S - - - - -	0	0	0/5
14448	- - - - H M D G - - - - V - - - - -	0	0	0/3
14496	- - - - H M D S - - - - V - - - - -	0	0	0/4
13724	- - - - H V T S - - - - S - - - - -	0	0	0/5
12896	- - - - H V P S - - - - A - - - - -	0	0	0/5
12898	- - - - H V P G - - - - A - - - - -	0	0	0/5
15954	- - - - L M D A - - - - S - - - - -	0	0	0/3
15956	- - - - L M P A - - - - S - - - - -	0	0	0/4
15958	- - - - L K D S - - - - S - - - - -	0	0	0/6
15960	- - - - L M D S - - - - S - - - - -	0	0	0/4
13728	- - - - N L D G - - - - S - - - - -	0	0	0/5
13452	- - - - L I D A - - - - S - - - - -	0	0	0/5
CONTROLS		0	0	0/50

GROUP
AGROUP
BGROUP
C



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Biochimie 84 (2003) 1181–1188

BIOCHIMIE

www.elsevier.com/locate/biochi

Protection against experimental *P. falciparum* malaria is associated with short AMA-1 peptide analogue α -helical structures

Marcia Cubillos^{a,1}, Luz Mary Salazar^{a,1}, Libardo Torres^b, Manuel Elkin Patarroyo^{a,b,*}

Polymerized peptide No.	Peptide Sequence	IFA ≥1:160 Post 2nd	IFA ≥1:160 Post 3rd	No. of protected monkeys	Group
4325	M I K S A F L P T G A F K A D R Y K S H	0	0	0/6	
13486	— — — A S — D — — — — — S P — — — — —	3(2560)	2(1280)	2/5	A
15516	— — — A — — H — — — — — M S — — W — — —	1(1280)	1(160)	1/4	A
20034	— — — A — — — — — — — — — — — — — — —	2(320)	1(160)	2/8	A
20032	— — — A — — — — — — — — — — — — — — —	1(160)	1(160)	1/8	A
22784	— — — V G — D — — — — — — — — — — — — — — —	1(2560)	ND	1/10	A
23404	— — — V S — D — — — — — M S — — — — — — — — —	2(320)	ND	1/10	A
23406	— — — V S — D D — — — — — M S — — — — — — — — —	1(320)	ND	1/10	A
14518	— — — V G — H — — — — — — — — — — — — — — —	1(320)	0	0/3	B
14520	— — — V G — H — — — — — M S P — — — — — — — — —	2(640)	0	0/3	B
17934	— — — A S — H V — — — — — M S — — — — — — — — —	2(320)	0	0/7	B
17936	— — — A S — H V — — — — — M S P — W — — — — —	2(640)	0	0/5	B
9192	— — — — — G — — — — — — — — — — — — — — —	0	0	0/5	C
9196	— —	0	0	0/5	C
10100	— — — R G — — — — — — — — — — — — — — —	0	0	0/5	C
10102	— — — F G — — — — — — — — — — — — — — —	0	0	0/5	C
10104	— — — G T — — — — — — — — — — — — — — —	0	0	0/5	C
12708	— —	0	0	ND	C
15512	— — — A — — — — — — — — — — — — — — —	0	0	0/4	C
15510	— — — A — — — — — — — — — — — — — — —	0	0	0/4	C
16000	— — — — — S — — — — — — — — — — — — — — —	0	0	0/5	C
16008	— — — — — — — — — — — H — — — — — P — — — — —	0	0	0/4	C
16002	— — — — — S — H — — — — — — — — — — — — — — —	0	0	0/5	C
16004	— — — — — S — H — — — — — — — — — — — — — — —	0	0	0/5	C
16006	— — — — — S — — — — — — — — — — — P — — — — —	0	0	0/5	C
17932	— — — A S — D — — — — — — — — — — — S P — W — — —	0	0	0/7	C
23178	— — — V S — D — — — — — — — — — — — — — — —	0	ND	0/9	C
23774	— — — V S — H D — — — — — M S — — — — — — — — —	0	ND	0/9	C
22424	— — — S S — H V — — — — — — — — — — — P — — — — —	0	ND	0/8	C
23000	— — — V — — D — — — — — — — — — — — S P — — — — —	0	ND	0/7	C
CONTROLS		0/60	0/60	0/60	

Plasmodium falciparum SERA protein peptide analogues having short helical regions induce protection against malaria

Marcia Cubillos^a, Martha Patricia Alba^a, Adriana Bermúdez^a,
Mary Trujillo^a, Manuel Elkin Patarroyo^{a,b,*}

Polymerized peptide No.	Peptide Sequence	IFA ≥1:320 Post 2nd	IFA ≥1:320 Post 3rd	No. of protected monkeys	Group
6737	D N I L V K M F K T N E N N D K S E L I	0/10	0/10	0/10	
12748	* - - - - - V I - - - - - *	1(2560)	1(2560)	1/4	A
13840	* - - - - - V I M - - - - - *	3(640)	1(320)	1/5	A
22834	* - - H - - - - V I - - - - - *	1(2560)	ND	2/9	A
22796	* - - - - - V I - - - - - *	1(5120)	ND	2/9	A
13844	* - - H - - - - V I M - - - - - *	1(320)	0/6	1/6	A
24096	* - - H - - - - V I M - - - - - *	1(5120)	ND	2/8	A
14094	* I - - - - - V I M - - - - - *	1(320)	2(320)	0/5	B
14096	* - - H - - R - V I M - - - - - *	3(640)	1(640)	0/4	B
24308	* - - H - - - - P M - - - - - *	1(640)	ND	0/8	B
9228	* G - - - - - - - - - - - *	0/5	0/5	0/5	C
9230	* - - G - - - - - - - - - *	0/5	0/5	0/5	C
9234	* - - - - - G - - - - - *	0/5	0/5	0/5	C
10154	* - - - - - M I - - - - - *	0/5	0/5	0/5	C
10152	* - - - - - M L - - - - - *	0/5	ND	0/5	C
10150	* - - - - - M F - - - - - *	0/5	0/5	0/5	C
16016	* - - - - - V D - - - - - *	0/5	0/5	0/5	C
17970	* - - H - - - - I - - - - *	0/8	0/7	0/7	C
22460	* - - - - - I - - - - - *	0/10	0/10	0/10	C
10148	* - - - - - R A - - - - - *	0/5	0/5	0/5	C
12754	* - - H - - - - V V I - - - - *	0/5	0/5	ND	C
23418	* - - H - - - - I M - - - - - *	0/10	ND	0/10	C
24392	* - - H - - - - A M - - - - - *	0/6	ND	0/6	C
CONTROLS		0/50	0/50	0/50	

Synthetic vaccines - Emerging principles

1. Plasmodia (falciparum & vivax) use CONSERVED as well as VARIABLE sequences to bind to host cells.
2. CONSERVED binding sequences are NOT antigenic, but VARIABLE sequences are
3. CONSERVED binding sequences are NOT immuno-genic in ANY of the tested species when administered as monomers, polymerized, associated with T helper epitopes or presented as MAPs. While VARIABLE sequences can be immunogenic they are strain-specific
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4. To overcome this problem the critical contact residues in the binding interaction have to be identified and changed
5. The critical contact residues have to be changed for aminoacids that ARE NOT commonly used during the mutational process in the variable sequences
6. **Physicochemical characteristics such as size, charge and volume have to be considered. Therefore A ↔ S; P ↔ D; F ↔ R; H ↔ L, K ↔ M; N ↔ I; Q ↔ E; C ↔ T; Y ↔ W; can be exchanged. G and C have special properties.**

NMR structure of *Plasmodium falciparum* malaria peptide correlates with protective immunity

Jindra Purmova, Luz Mary Salazar, Fabiola Espejo, Mary Helena Torres, Marcia Cubillos, Elizabeth Torres, Yolanda Lopez, Raul Rodriguez, Manuel Elkin Patarroyo*

Fundación Instituto de Immunología de Colombia (FIDIC), Carrera 50, No. 26-00, Bogotá, Colombia

Received 6 November 2001; received in revised form 12 February 2002; accepted 26 March 2002

FEBS 26417

FEBS Letters 527 (2002) 95–100

Protection against experimental malaria associated with AMA-1 peptide analogue structures

Luz M. Salazar, Martha P. Alba, Maria H. Torres, Martha Pinto, Ximena Cortes, Libardo Torres, Manuel E. Patarroyo*

2250

J. Med. Chem. 2003, 46, 2250–2253

Distorting Malaria Peptide Backbone Structure to Enable Fitting into MHC Class II Molecules Renders Modified Peptides Immunogenic and Protective

Gladys Cifuentes,[†] Manuel Elkin Patarroyo,^{*,†,‡} Mauricio Urquiza,^{†,‡} Luis E. Ramirez,[†] Claudia Reyes,[†] and Raul Rodriguez[†]

6746 SERA peptide analogues immunogenicity and protective efficacy against malaria is associated with short α helix formation: Malaria protection associated with peptides α helix shortening

Martha Patricia Alba^a, Luz Mary Salazar^a, Alvaro Puentes^a, Martha Pinto^a, Elizabeth Torres^a, Manuel Elkin Patarroyo^{a,b,*}

Shortening and modifying the 1513 MSP-1 peptide's α -helical region induces protection against malaria

Fabiola Espejo,¹ Adriana Bermúdez,¹ Elizabeth Torres, Mauricio Urquiza, Raúl Rodríguez, Yolanda López, and Manuel Elkin Patarroyo^{*}

Biol. Chem., Vol. 384, pp. 000–000, October/November 2003 · Copyright © by Walter de Gruyter · Berlin · New York

Adriana Bermúdez¹, Gladys Cifuentes¹, Fanny Guzmán¹, Luz Mary Salazar¹ and Manuel Elkin Patarroyo^{1,2,*}
Immunogenicity and Protectivity of *Plasmodium falciparum* EBA-175 Peptide and Its Analog Is Associated with α -Helical Region Shortening and Displacement

Eur. J. Biochem. 270, 1–7 (2003) © FEBS 2003

doi:10.1046/j.1432-1033.2003.03780.x

Modified merozoite surface protein-1 peptides with short alpha helical regions are associated with inducing protection against malaria

Mary H. Torres^{1*}, Luz M. Salazar^{1*}, Magnolia Vanegas¹, Fanny Guzman¹, Raul Rodriguez¹, Yolanda Silva¹, Jaiver Rosas¹ and Manuel E. Patarroyo^{1,2}

PROTEINS:Structure,Function,and Genetics 50:400 –409 (2003)

Alpha Helix Shortening in 1522 MSP-1 Conserved Peptide Analogs is Associated with Immunogenicity and Protection Against *P. falciparum* Malaria

Marcia Cubillos,¹ Fabiola Espejo,¹ Jindra Purmova,¹ Juan C. Martinez,² and M.E. Patarroyo^{1,2}

Analysis of a *Plasmodium falciparum* EBA-175 peptide with high binding capacity to erythrocytes and their analogues using ¹H NMR

Gladys Cifuentes, Fanny Guzmán, Martha Patricia Alba, Luz Mary Salazar, and Manuel Elkin Patarroyo*

Induction and displacement of an α helix in the 6725 SERA peptide analogue confers protection against *P. falciparum* malaria

Martha Patricia Alba^a, Luz Mary Salazar^a, Jindra Purmova^a, Magnolia Vanegas^a, Raul Rodriguez^a, Manuel Elkin Patarrovo^{a,b,*}

¹H-NMR structures of the *Plasmodium falciparum* 1758 erythrocyte binding peptide analogues and protection against malaria

Fanny Guzman^{a,1}, Katherine Jaramillo^{a,1}, Luz M. Salazar^a, Angela Torres^a, Augusto Rivera^b, Manuel E. Patarrovo^{a,b,*}

Orientating Peptide Residues and Increasing the Distance between Pockets to Enable Fitting into MHC–TCR Complex Determine Protection against Malaria

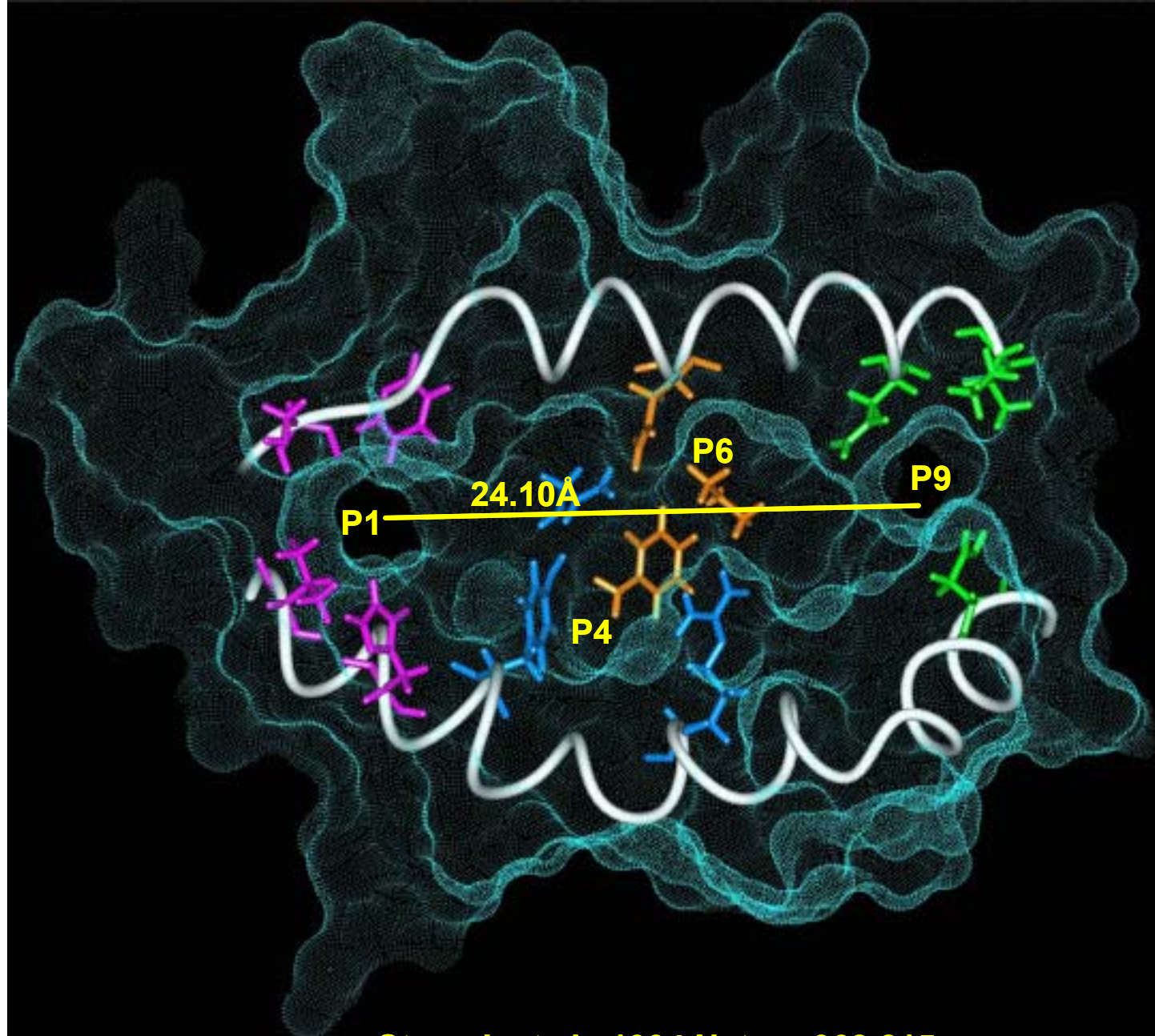
Gladys Cifuentes,[§] Fabiola Espejo,[§] Luis Eduardo Vargas,[§] Carlos Parra,^{§,‡} Magnolia Vanegas,[§] and Manuel Elkin Patarrovo^{*,§,‡}

Received February 10, 2004; Revised Manuscript Received March 24, 2004



The MHC molecules
the peptide and
the T cell receptor
complex

Class-II Molecule

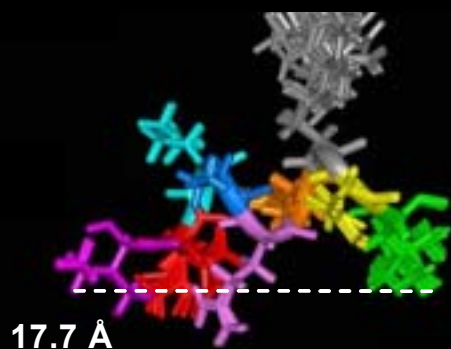


Stern J. et al., 1994 Nature 368,215

Native

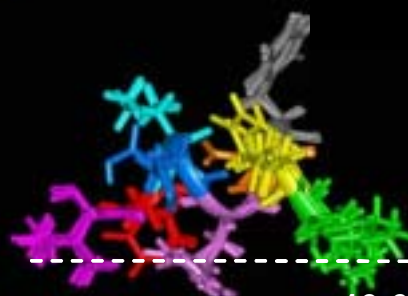
Protective

1815



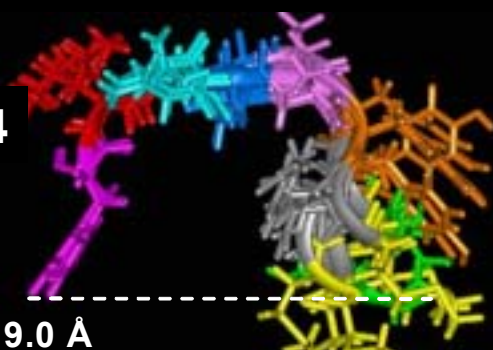
17.7 Å

24292



19.6 Å

4044



19.0 Å

10008



22.2 Å

1522



16.9 Å

13446



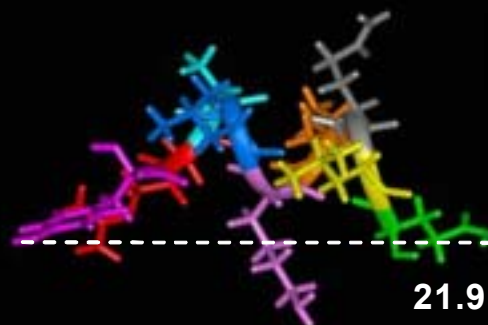
20.2 Å

1585



14.3 Å

11860



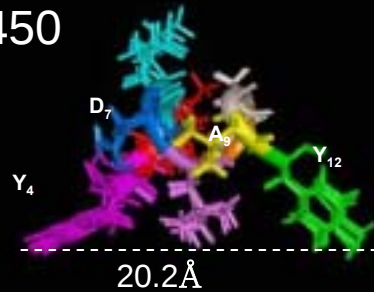
21.9 Å

Patarroyo M. E. et al. 2004. *ChemBioChem*. 5: 1588

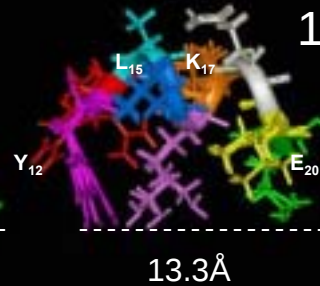
Immunogenic and Protection inducers

Short-lived antibodies inducers

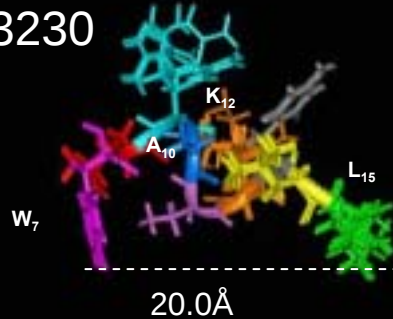
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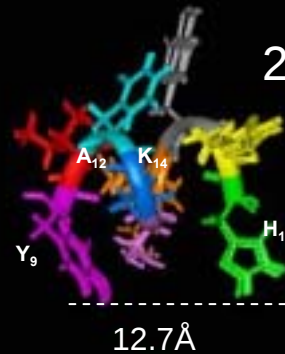
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23230



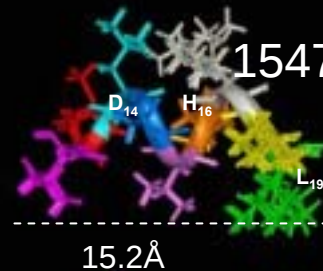
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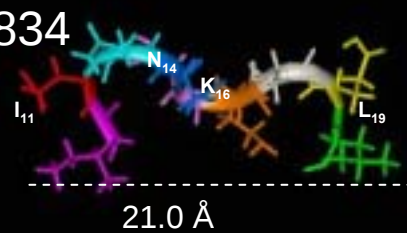
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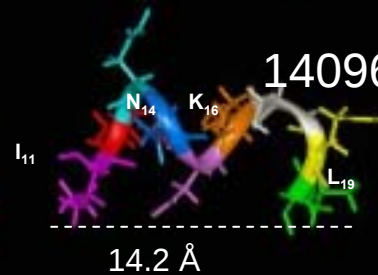
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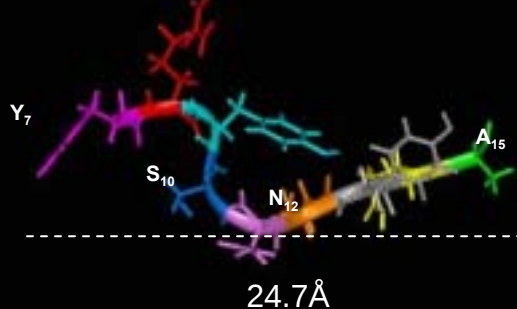
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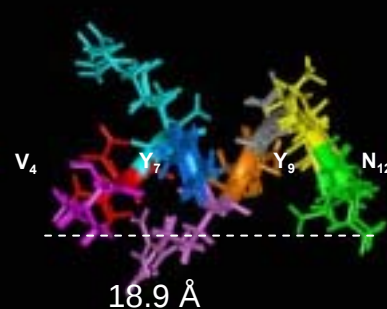
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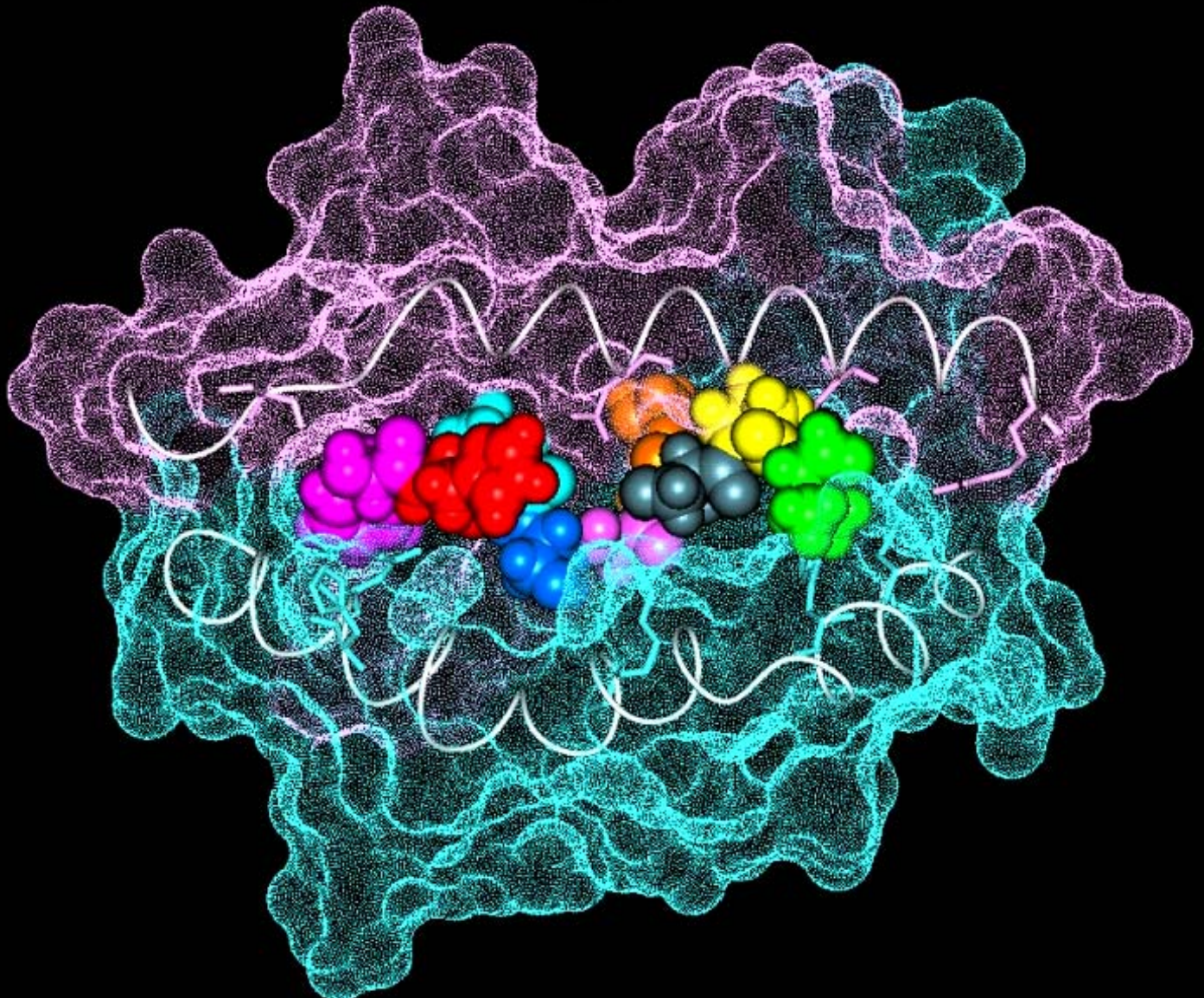
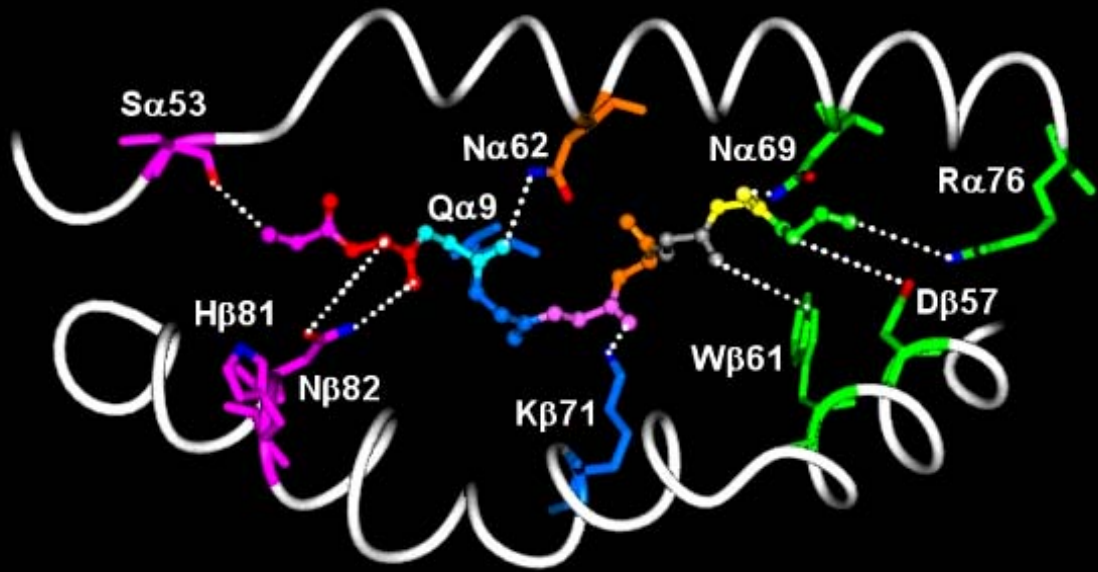
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10000



HLA-DRB1*0401 and 10022 Peptide

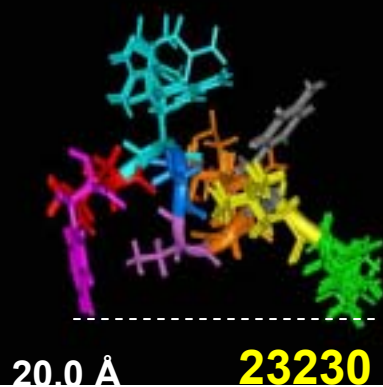
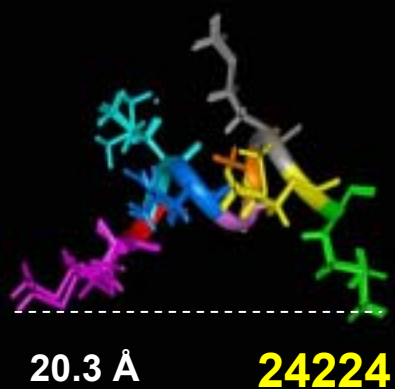
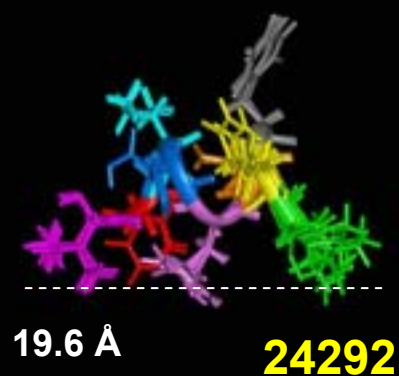
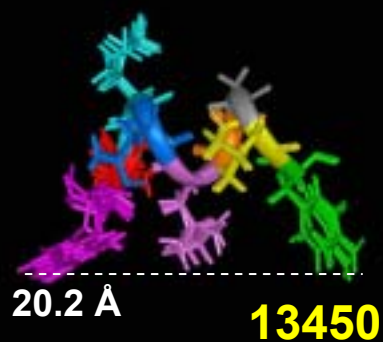


Salazar L. M. et al. 2004. *Biochem. Biophys. R. Comun* . 322:119

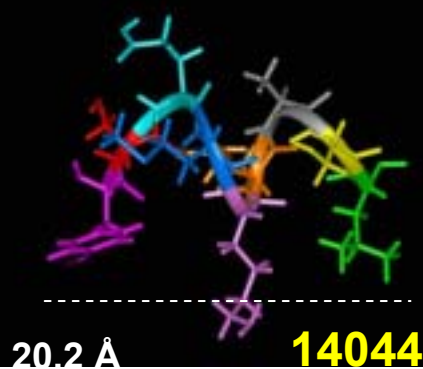
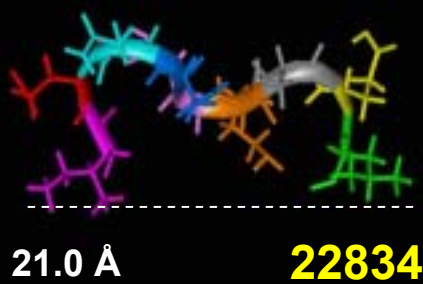
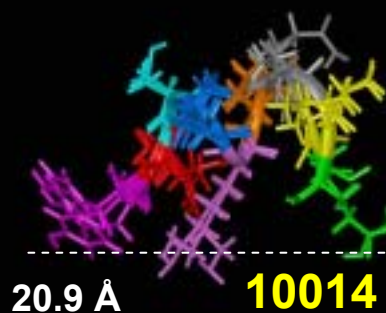
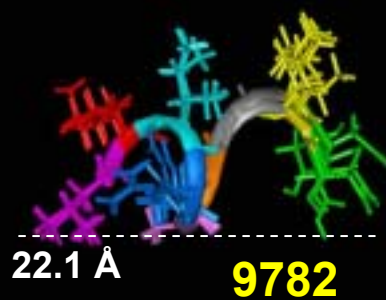
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6. Physicochemical characteristics such as size, charge and volume have to be considered. Therefore $A \leftrightarrow S$; $P \leftrightarrow D$; $F \leftrightarrow R$; $H \leftrightarrow L$, $K \leftrightarrow M$; $N \leftrightarrow I$; $Q \leftrightarrow E$; $C \leftrightarrow T$; $Y \leftrightarrow W$; can be exchanged. G and C have special properties.
7. There must be a $23.5 \text{ \AA} \pm 3.5 \text{ \AA}$ distance between those residues fitting into Pockets 1-9 of Class II molecules.

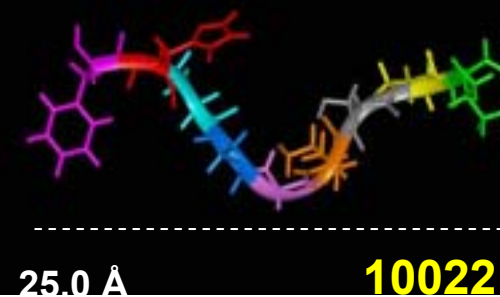
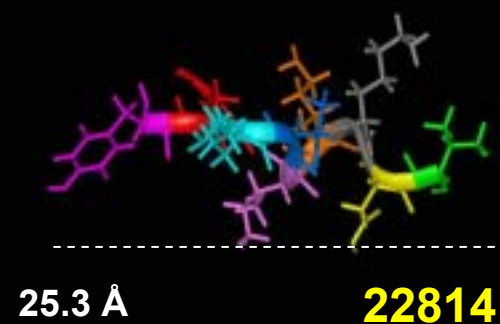
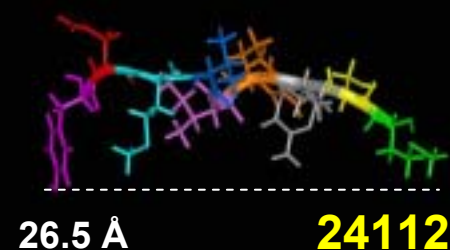
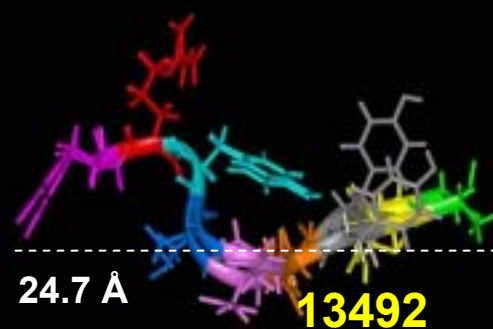
DR β 1*0301



DR β 1*1101



DR β 1*0401



DRβ1*0301



13450

DRβ1*1101



9782

DRβ1*0401



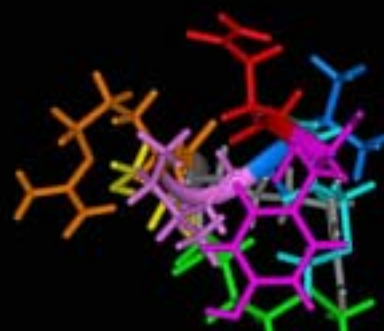
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24292



10014



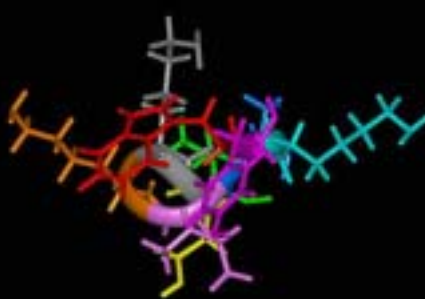
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24224



22834



22814



23230

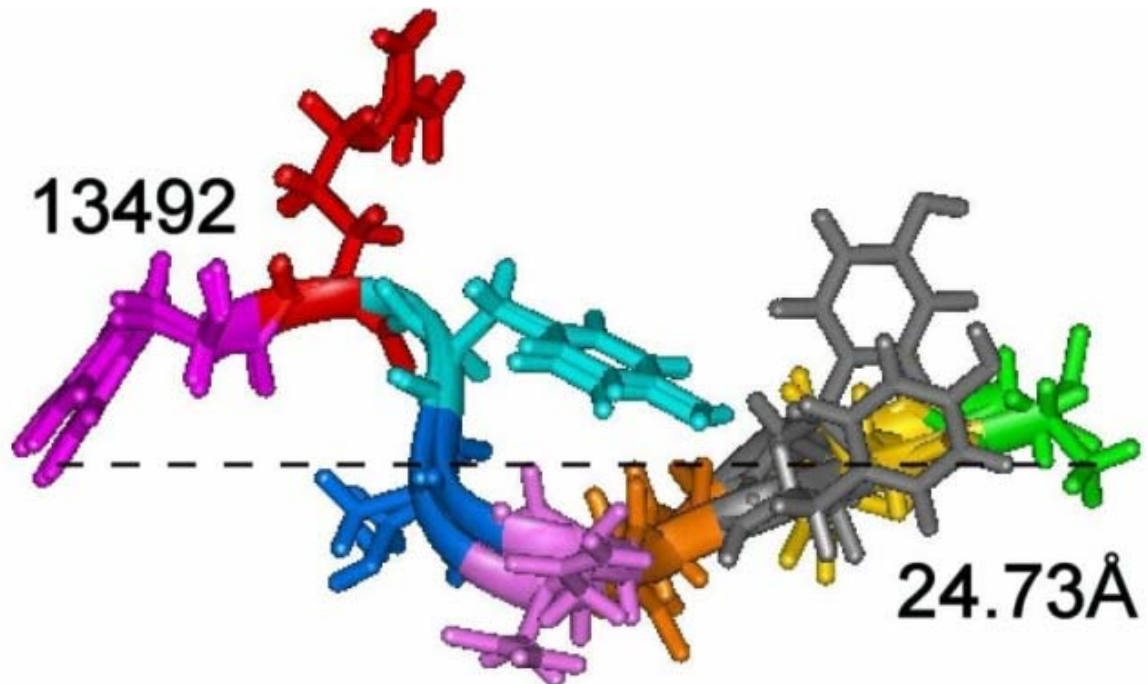


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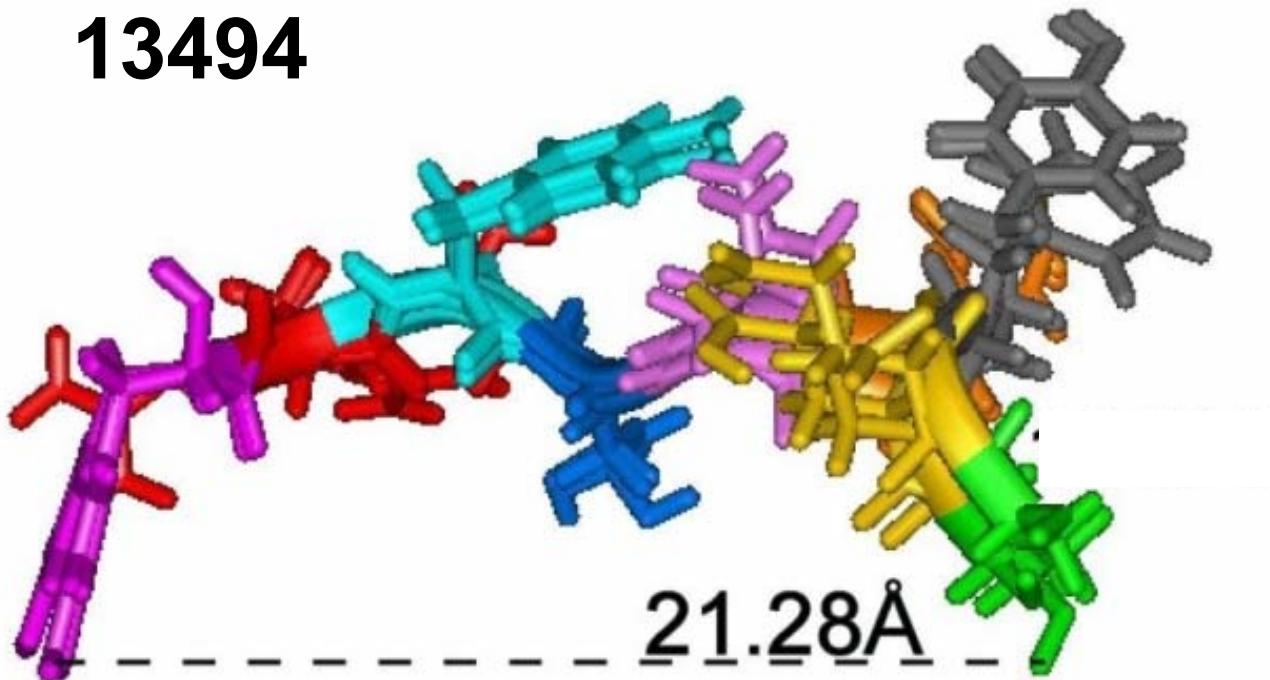


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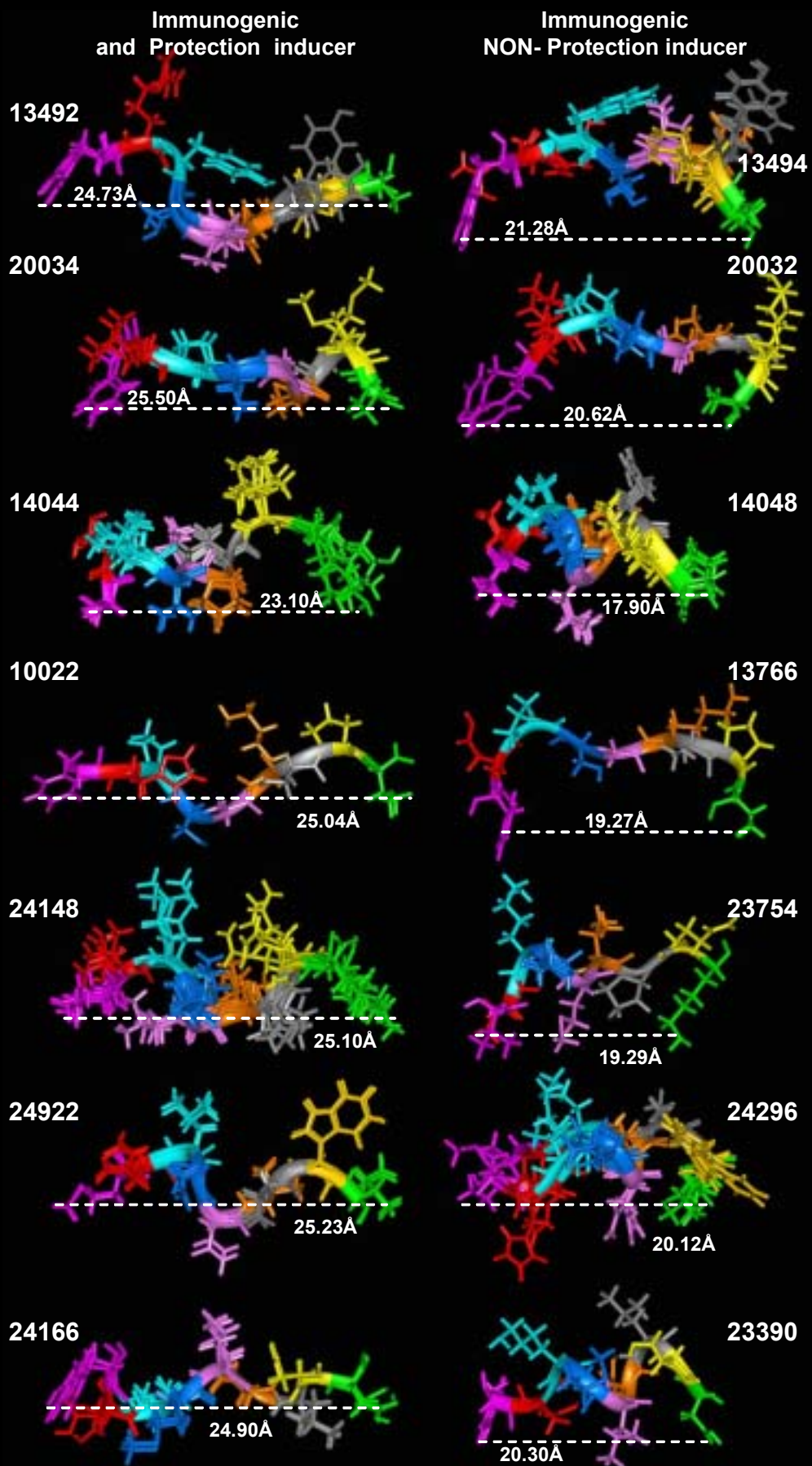
Immunogenic and Protection Inducer



Immunogenic and NON-Protection Inducer



Patarroyo M. E. et al. 2004, Submitted to *Science*



Synthetic vaccines - Emerging principles

1. Plasmodia (falciparum & vivax) use CONSERVED as well as VARIABLE sequences to bind to host cells.
2. CONSERVED binding sequences are NOT antigenic, but VARIABLE sequences are
3. CONSERVED binding sequences are NOT immuno-genic in ANY of the tested species when administered as monomers, polymerized, associated with T helper epitopes or presented as MAPs. While VARIABLE sequences can be immunogenic they are strain-specific
4. To overcome this problem the critical contact residues in the binding interaction have to be identified and changed
5. The critical contact residues have to be changed for aminoacids that ARE NOT commonly used during the mutational process in the variable sequences
6. Physicochemical characteristics such as size, charge and volume have to be considered. Therefore $A \leftrightarrow S$; $P \leftrightarrow D$; $F \leftrightarrow R$; $H \leftrightarrow L$, $K \leftrightarrow M$; $N \leftrightarrow I$; $Q \leftrightarrow E$; $C \leftrightarrow T$; $Y \leftrightarrow W$; can be exchanged. G and C have special properties.
7. There must be a $23.5 \text{ \AA} + 3.5 \text{ \AA}$ distance between those residues fitting into Pockets 1-9 of Class II molecules.
8. All the changes made to the CONSERVED binding peptides are designed to achieve a perfect fit with the MHC molecules and the TCR



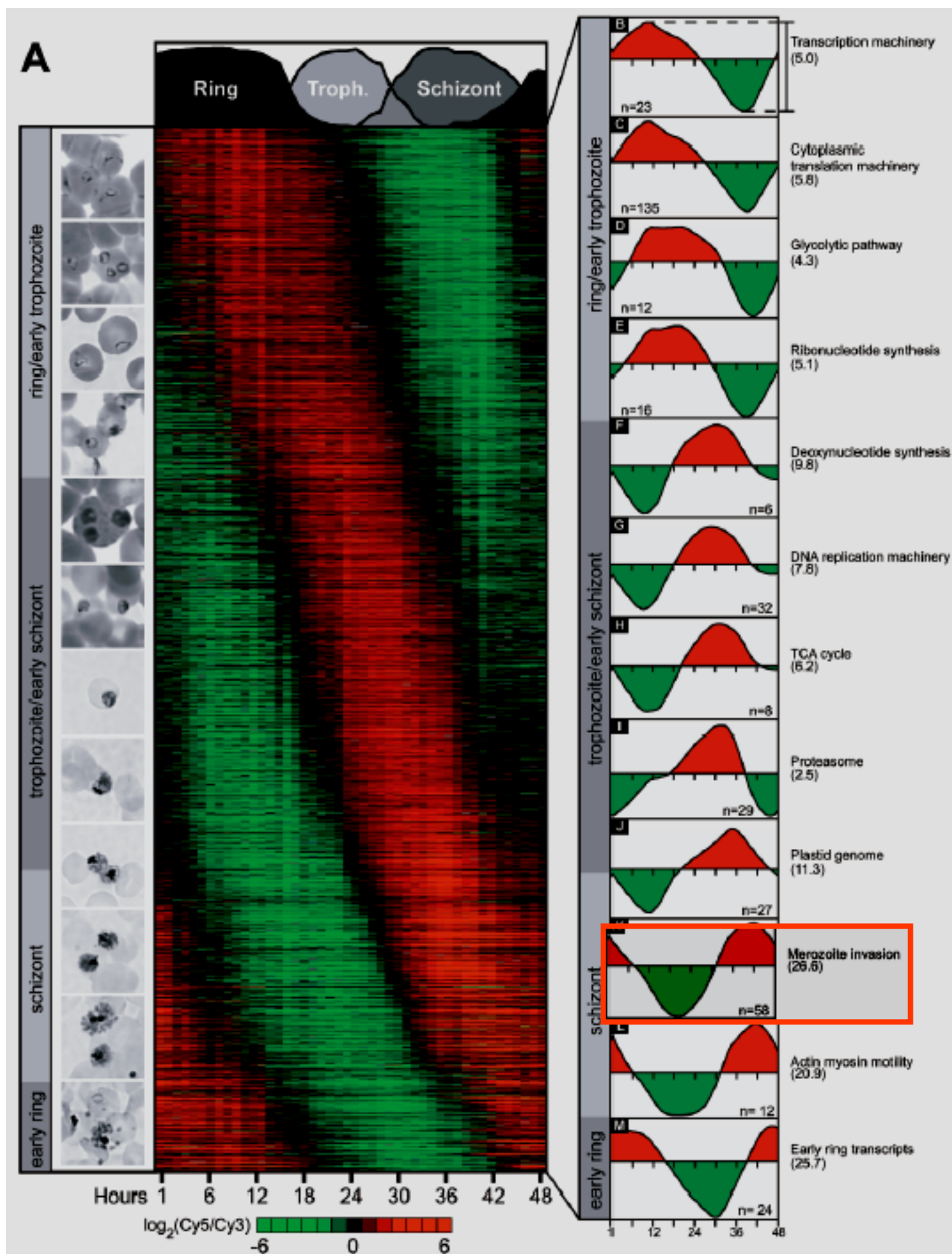
Reinherz E. L. et al., 1999 Science 286, 1973

Synthetic vaccines - Emerging principles

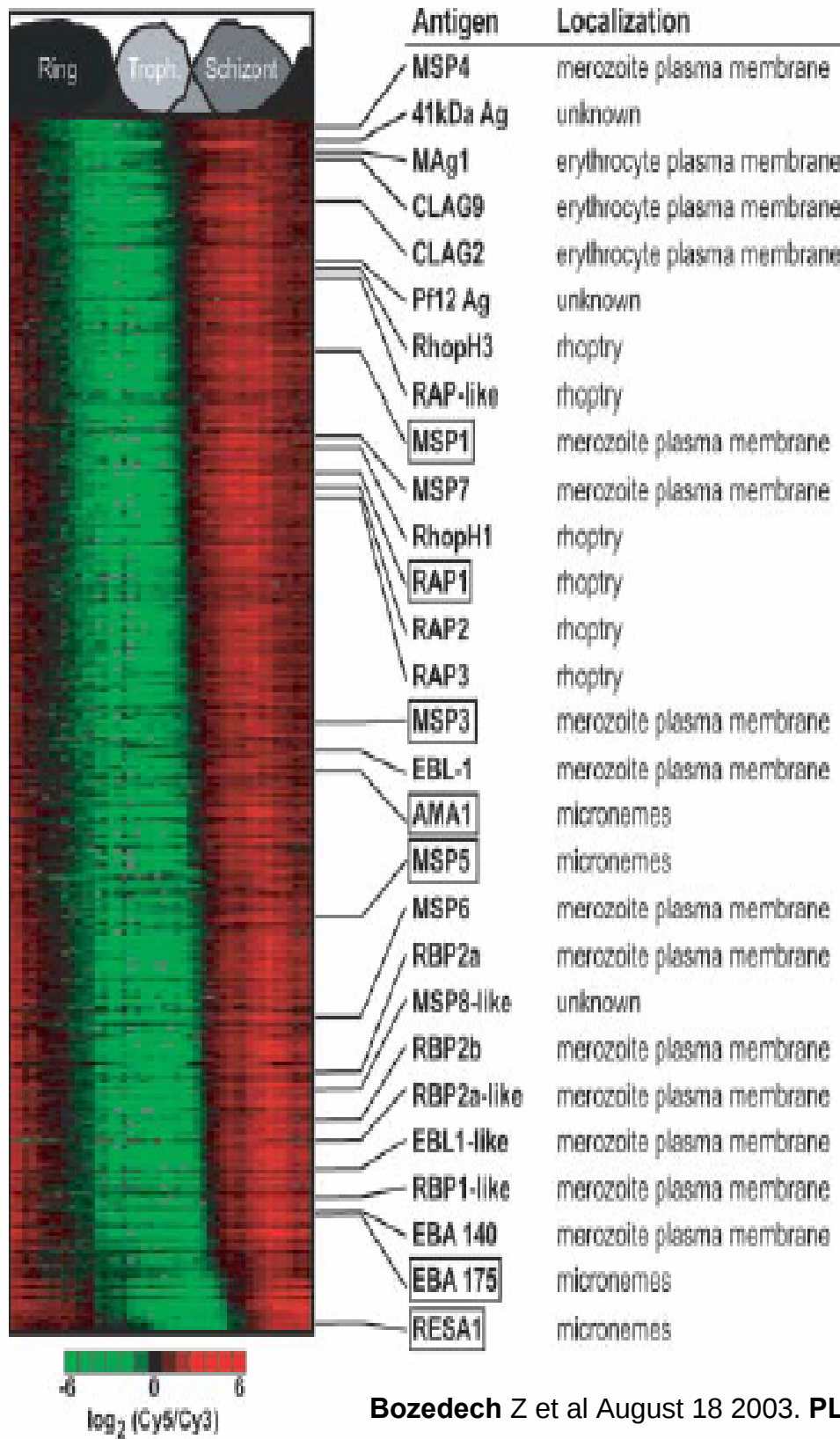
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8. All the changes made to the CONSERVED binding peptides are designed to achieve a perfect fit with the MHC molecules and the TCR
- 9 **Structural considerations together with affinity, biological and Immunological activities should guide the design of synthetic peptide vaccines.**

COLFAVAC®

PEP	SEQUENCE	HLA-DR
24148	MLNI SMLQTVMMTPQK	DR1
24150	WKSYSVD DNI PMNMSLIHKH	DR3
22812	NNDRIYDMNHLMIKMHILAI	DR4W4
22814	NDKLYRMEYWKTIKKDVW	DR12
24292	LTNQNINIDQEFNLMKHGFH	DR3
24166	FNNI PSRYNLYDKMLPLDD	DR11
23394	KKMNMMENVIIYIMKNQNLFK	DR12
24112	SKYSNTFNINAYNMVIRRS	DR12
22782	GEDAEVAGTQWFVPSGKVPVFG	DR1
23406	MIKVSFDDTGAFMSDRYKSH	DR3
22822	HTTYSEEKRASHTTPVLKEK	DR11
24300	ATADYFRHLVTPQNLE	DR15
22720	TDVNRVRYVNNYESNPHK	DR4W10
23422	KETNNAISFMSNAGSLEKK	DR4W10
24096	DNIHVKMFKVIMNNDKSELI	DR12
24216	DQGNTITAWNRAAKFHLETI	DR4W14
23426	KKVQNLTGDDTADLATNIVG	DR3
24224	KSKKHMDLDGEMMAKKLKD	DR3
24228	SAFDDNLTAMNAMGLILNKR	DR1
24238	VAFNRFHVGTHPAPGKTNLY	DR11
24244	VWDEWSPVSTAVAMGTRSRK	DR1
24246	SPTS SVTVGKGAFSPKRE	DR1
24254	GAATPYSGEPSPFDEVLGEE	DR4W15
24318	HNMPNDDIRPVDEPANAN	DR4W13



PHASEOGRAM OF PUTATIVE VACCINE TARGETS



Bozdech Z et al August 18 2003. **PLOS Biology** 1, 001

High RBC binding activity peptides



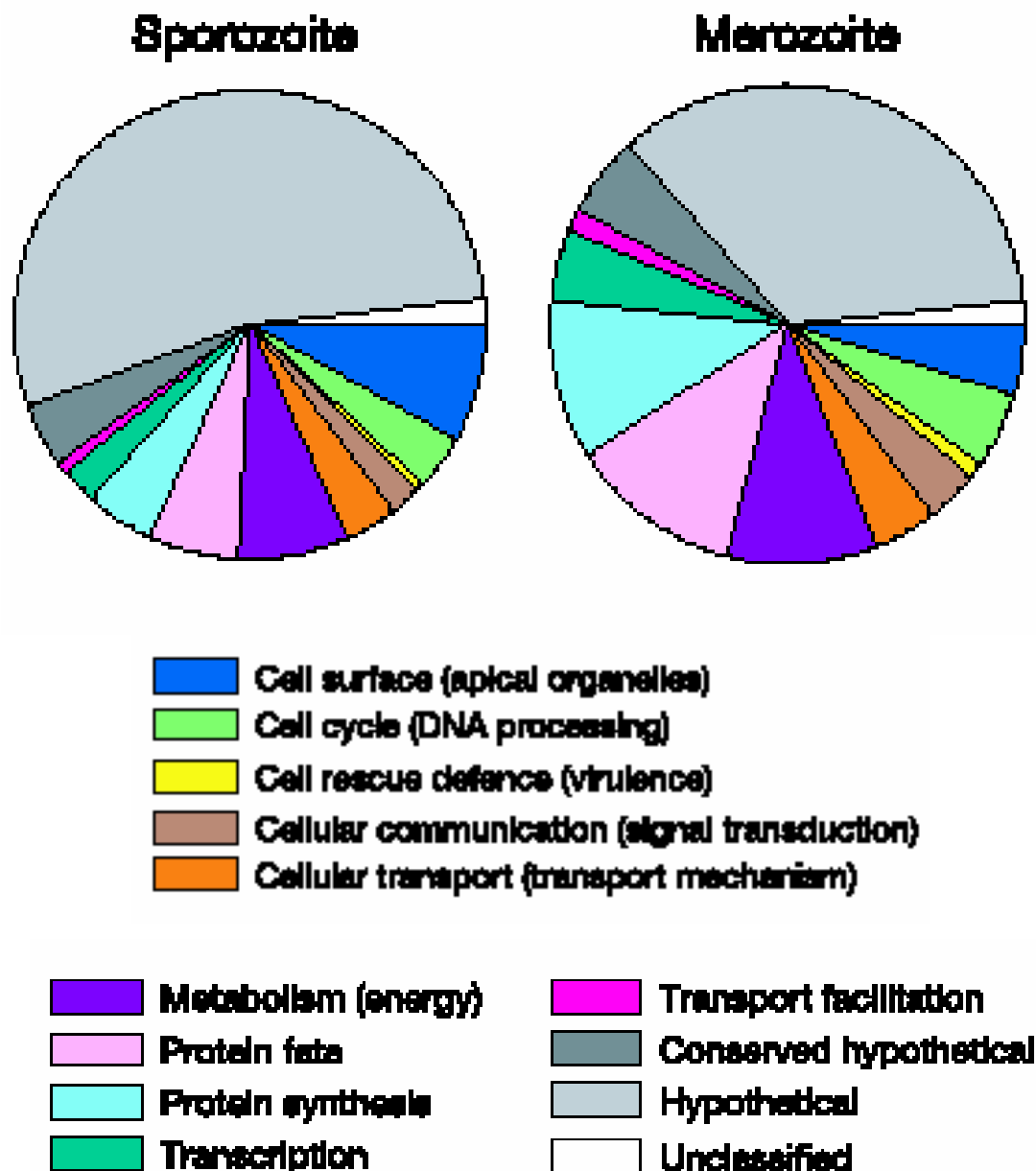
PROTEIN	C	V	Reference
MSP1	3	6	<u>Urquiza M. et. al . 1996. <i>Parasite Immunology</i> . 18: 515-526</u>
RESA	2	–	<u>Vera R. et al . 2000. <i>Vaccine</i>. 18: 1289-1293</u>
SERA	6	1	<u>Puentes A. et. al . 2000. <i>Parasitology International</i>. 49: 105-117</u>
MSA2	1	2	<u>Ocampo M. et. al . <i>J. Peptide Research</i>. 55: 216-233</u>
EBA175	6	–	<u>Rodríguez L.E et. al. 2000. <i>Parasitology</i>. 120: 225-235</u>
GBP130	1	–	<u>Suárez J. et. al . 2000. <i>Mem. Inst. Oswaldo Cruz</i> . 95: 495-501</u>
HRP I, II, II	3	1	<u>López R. et a l. 2000. <i>Acta Tropica</i>. 75: 349-359</u>
ABRA	5	–	<u>Curtidor H. et. al . 2001. <i>Vaccine</i>. 19: 4496-4504</u>
AMA1	4	3	<u>Urquiza M. et. al . 2001. <i>Vaccine</i>. 19: 508-513</u>
EBA140	6	–	<u>Rodríguez L.E.. et. al . 2003. <i>J. Peptide Research</i>. 62: 175-184</u>
NBP1	2	–	<u>Valbuena J. et al . 2003. <i>Peptides</i> . 24: 1007-1014</u>
MSP8	5	–	<u>Puentes A. et. al . 2003. <i>Peptides</i> .24: 1015-1023</u>
MAEBL	9	–	<u>Ocampo M. et al . 2004. <i>Biochem. Biophys. Res. Comm</i>. 315: 319-329</u>
RAP2	4	–	<u>López R. et al . 2004. <i>Biochemie</i> . 86: 1-6</u>
RBP2 Ha/Hb	9	3	<u>Ocampo M. et. al . 2004. <i>Parasitol. Int</i>. 53: 77-88</u>
RAP1	4	–	<u>Curtidor H. et. al . 2004. <i>Vaccine</i> . 22: 1054-1062</u>
EBA160	5	–	<u>Valbuena J. et al . 2004. <i>Biochem. Biophys. Res. Comm</i> . 321: 835-844</u>
CLAG3	5	–	<u>Ocampo M. et. al . 2005. <i>Proteins Science</i> . 14: 504-513</u>
EBL1	5	–	<u>Curtidor H. et. al . 2005. <i>Protein Science</i> . 14: 464-473</u>
JESEBL	5	–	<u>Vera R. et al . 2004. <i>Biochemie. In press</i></u>
STEVOR	–	3	<u>García J. et al . 2004. <i>Peptides. In press</i></u>
MSP10	3	–	<u>Puentes A. et al . 2004. <i>Biochemie. In press</i></u>
MSP3	3	–	<u>Rodríguez L.E. et. al . 2004. <i>Submitted to Protein Science</i></u>
SORTILIN	6	–	<u>Vera R. et al . 2004. <i>Submitted to Biochemistry</i></u>
RESA-LIKE	3	–	<u>Rodríguez L.E. et. al . 2004. <i>Submitted to J. Int. Parasitol.</i></u>
TryThrA	4	–	<u>Curtidor H. et. al . 2004. <i>Submitted to Chembiochem</i></u>
HAP	2	–	<u>Valbuena J. et. al . 2004. <i>Submitted to Biol. Chem.</i></u>
MSP6	2	1	<u>López R. et al . 2004. <i>Submitted to Peptides</i></u>

C : CONSERVED

V: VARIABLE



Functional profile of proteins expressed in different *P. falciparum* life-cycle stages



Plasmodium sporozoite surface and/or apical organelle proteome

Protein	Other stages	Subcellular Localization	Conserved pep
Cs	-	Surface	2
TRAP	-	Surface, micronemes	7
STARP	RBC, LS	Surface	5
SALSA	RBC, LS	Surface	2
LSA3	LS	Surface	11
LSA1	LS		1
EBA175	Mz	Surface, micronemes	
AMA1	Mz	Surface, micronemes	
MAEBL	-	Surface, micronemes	
MCP1	MZ	?	
PEMP3	RBC, LS	Surface	
SPATR	RBC	Surface	
PfEMP1	ïRBC		
STEVOR	ïRBC		
RIFINS	ïRBC		
P52	-	?	
P235	Mz		
PPLP1	-	Micronemes	
SPECT	-	Micronemes	

Ramsés López et al 2001, J. Pept. Res, 58: 285

Control	PI GEEPDHQL DPAFGANSNNPDWDENP		
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Plasmodium falciparum circumsporozoite (CS) protein peptides specifically bind to HepG2 cells

Jorge E. Suarez *, Mauricio Urquiza, Alvaro Puentes, Javier E. Garcia, Hernando Curtidor, Marisol Ocampo, Ramses Lopez, Luis E. Rodriguez, Ricardo Vera, Marcia Cubillos, Maria H. Torres, Manuel E. Patarroyo

PEPTIDE NUMBER	Sequence	BINDING ACTIVITY
4370	1 MMRKLAILSVSSFLFVEALFQEY 22	
4371	8 LSVSSFLFVEALFQEYQCYG 27	
4372	13 FLFVEALFQEYQCYGSSSNT 32	
4373	18 ALFQEYQCYGSSSNTLRLNE 37	
4374	23 YQCYGSSSNTLRLNELNYDN 42	
4375	28 SSSNTLRLNELNYDNAGTNL 47	
4376	33 RVLNELNYDNAGTNLYNELE 52	
4377	38 LNYDNAGTNLYNELEMNYYG 57	
4378	43 AGTNLY ELEMNYYGKQENW 62	
4379	48 YNELEMNYYGKQENWYSLKK 67	
4380	53 MNYYGKQENWYSLKKNSRSL 72	
4381	58 KQENWYSLKKNSRSLGENDD 77	
4382	63 YSLKKNSRSLGENDDGNNED 82	
4383	68 NSRSLGENDDGNNEDNEKLRY 87	
4384	73 GENDDGNNEDNEKLKPKHKY 92	
4525	78 GNNEDNEKLKPKHKKLKQPY 97	
4386	83 NEKLKPKHKKLKQPADGNPY 102	
4592	103 DPANPVDPNANPNVY 112	
4593	113 NANPNANPNANPY 122	
4387	281 NKNNQGNGQGHNPNDPNRNY 300	
4388	286 GNGQGHNPNDPNRNVDENAY 305	
4389	291 HNMPNDPNRNVDENANANSAY 310	
4390	296 DPNRNVDENANANSVKNNNY 315	
4391	301 VDENANANSVKNNNNEEPSY 320	
4392	306 NANSVKNNNNEEPSDKHIKY 325	
4393	311 VKNNNNEEPSDKHIKEYLNK 330	
4394	316 NEEPSDKHIKEYLNKIQNSL 335	
4395	321 DKHIKEYLNKIQNSLSTEWS 340	
4396	326 EYLNKIQNSLSTEWSPCSVT 345	
4397	331 IQNSLSTEWSPCSVTCGNGIY 350	
4398	336 STEWSPCSVTCGNGIQVRIKY 355	
4399	341 PCSVTCGNGIQVRIKPGSANY 360	
4400	346 CGNGIQVRIKPGSANKPKDEY 365	
4401	351 QVRIKPGSANKPKDELIDYAN 370	
4402	356 PGSANKPKDELIDYANDIEKK 375	
4403	361 KPKDELIDYANDIEKKICKMEKCS 380	
4404	366 LDYANDIEKKICKMEKCSSV 385	
4405	371 DIEKKICKMEKCSSVFNVVNY 390	
4406	376 ICKMEKCSSVFNVVNSSIGLY 395	
4407	381 KCSSVFNVVNSSIGLIMVLSY 400	
4408	386 FNVVNSSIGLIMVLSFLFN 405	

83

RTS-S

383

Polimorphic sequence found in different *P. falciparum* isolates **Corresponding to amino acids 326 to 345** **Of the NF54 *P. falciparum***

Moreno A. et al. 1993 **J Immunol.** 151:489.

Isolates	Origin	Sequence												
NF54,XP _{8,9} , X _{10,11} ,427 _{1-4,6-10} 3D7	Wets Africa, Honduras, The Netherlands	<u>E</u>	Y	L	<u>N</u>	<u>K</u>	I	<u>Q</u>	N	S	<u>L</u>	S	T	E
ItG2G1	Brazil	K			K									
MS2,SO6	Thailand, Brazil	K			K	R								
LE5, 366 ₉	Liberia, Gambia	K			K	T								
366 ₁ , 399 ₁₋₁₀ , 406 ₁₋₉ , 419 ₁₀	Gambia	K			K	T	K							
366 _{8,10}	Gambia	K			Q									
406 ₁₀ , 419 ₁₋₉	Gambia	K			Q		K							
366 ₂₋₇	Gambia	K			Q		R							
Xs,XP _{12,13} HB3, D10, T4, T9-101,	Honduras, The Netherlands, Papua New Guinea,	Q			K									
WEL, B11, PA, FCR3	Thailand, West Africa, Uganda, Gambia													
S24, T9-98	Angola, Thailand	Q			K	T	K							
7G8	Brazil	Q			K		K				I			

^aAmino acids numbering is based on the CS protein of the NF54 strain of *P. falciparum* . The polimorphic residues are underline, and only amino acids that differ from the NF54 sequence are shown for each strain or isolate. The subscript numbers correspond to the isolate name described in materials and methods

SOME OTHER

VACCINES

IN PROGRESS

AT FIDIC

Plasmodium vivax Duffy binding protein peptides specifically bind to reticulocytes

Marisol Ocampo*, Ricardo Vera, Luis Eduardo Rodriguez, Hernando Curtidor, Mauricio Urquiza, Jorge Suarez, Javier Garcia, Alvaro Puentes, Ramsés Lopez, Mary Trujillo, Elizabeth Torres, Manuel Elkin Patarroyo

Peptide Number	Sequence		Reticulocytes		Erythrocytes		
			2%	4%	2%	4%	
1616	16 SQVNNVLLERTIETLLECKN	97					I
1617	36 EYVKGNGYKLAQHHCVEE	67					
1618	66 DNLERWLQGTNERRSEENIK	77					
1619	76 YKYGVTELKIKYAQMNGKRS	97					
1620	96 SRILKESIYGAHNFGGNSYM	117					
1621	116 EGKDGDDKTGEEKDGEHKT	137					II
1622	136 SKTDNGKGANNLVMLDYETS	157					
1623	156 SNGQPAGTLDNVLEFVTGHE	177					
1624	176 GNSRKNSNGGNPYDIDHKK	197					
1625	196 TISSAIINHAFLQNTVMKNC	217					
1626	216 NYKRKRERRERDWCNTKKDVC	237					
1627	236 IPDRRYQLCMKELTNLVNNT	257					
1628	256 DTNFHRDITFRKLYLKRKLI	277					
1629	276 YDAAVEGDLLLKLNNYRYNK	297					
1630	296 DFCKDIRWSLGDFGDIIMGT	317					
1631	316 DMEGIGYSKVVENNLRISIFG	337					
1632	336 TDEKAQQRKQWWNESKAQI	357					
1633	356 WTAMMYSVKKRLKGNFIWIC	377					
1634	376 KLNVAVNIEPQIYRWIREWG	397					
1635	396 RDYVSELPTEVQKLKEKCDG	417					
1636	416 KINYTDKKVCKVPPCQNAACK	437					
1637	436 SYDQWITRKKNQWDVLSNKF	457					
1638	456 ISVKNAEKVQTAGIVTPYDI	477					
1639	476 LKQELDEFNEVAFENEINKR	497					III-V
1640	496 DGAYIELCVCVVEEAKKNTQ	517					
1641	516 EVVTNVDNAAKSQATNSNPI	537					
1642	536 SQPVDSSKAKEKVPDSTHGN	557					
1643	556 VNSGQDSSTTGKAVTGDGQN	577					
1644	576 GNQTPAESDVQRSDIAESVS	597					
1645	596 AKNVDPQKSVSKRSDDTASV	617					
1646	616 TGIAEAGKENYGASNSRPSE	637					
1647	636 STVEANSPGDDTVNSASIPV	657					
1648	656 VSGENPLVTPYNGLRHSDKDN	677					
1649	676 SDSGPAESMANPDSNSKGE	697					VI
1650	696 TGKGQDNDMAKATKDSNSSS	717					
1651	716 DGTSSATGDTTDAVDREINK	737					
1652	736 GVPEDRDKTGVGSKDGGGEDN	757					
1653	756 SANKDAATVVGEDRIRENSA	777					
1654	776 GGSTNDRSKNDTEKNGASTP	797					
1655	796 DSKQSEDATLSKTESLEST	817					
1656	816 ESGDRTTNDTTNSLENKNGG	837					
1657	836 KEKDLQKHDFKSNDDTPNEEP	857					
1658	856 NSDQTTDAEGHDRDSIKNDK	877					
1659	876 AERRKHMNKDTFTKNTNSHH	897					transm
1660	896 LNSNNNLSNGKLDIKEYKYR	917					
1661	916 DVKATREDIILMSSVRKCNN	937					
1662	936 NISLEYCNSVEDKISSNTCS	957					
1663	956 REKSKNLCCSISDFCLNYFD	977					
1664	976 VYSYEYLSCKMKKEFEDPSYK	997					
1665	996 CFTKGGFKDKTYFAAAGALL	1017					
1666	1016 ILLLLIASRKMKNDSSEAT	1037					
1667	1036 FNEFEYCDNIHRIPLMPNN	1057					
1668	1056 IEHMQPSTPLDYS	1070					



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Vaccine

Vaccine 21 (2003) 4133–4144

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A highly infective *Plasmodium vivax* strain adapted to *Aotus* monkeys Quantitative haematological and molecular determinations useful for *P. vivax* malaria vaccine development

Yago Pico de Coaña^a, Josefa Rodríguez^a, Eduar Guerrero^a, Carlos Barrero^a,
Raúl Rodríguez^a, Marcela Mendoza^b, Manuel Alfonso Patarroyo^{a,*}



Experimental Parasitology 100 (2002) 131–134

Experimental
Parasitology

www.academicpress.com

Research Brief

Plasmodium vivax: parasitemia determination by real-time quantitative PCR in *Aotus* monkeys

Juan Carlos Polanco, Josefa Antonia Rodríguez, Vladimir Corredor,
and Manuel Alfonso Patarroyo*



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Vaccine

Vaccine 21 (2003) 4133–4144

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Review

Splenectomised and spleen intact *Aotus* monkeys' immune response to *Plasmodium vivax* MSP-1 protein fragments and their high activity binding peptides

Adriana Yanett Sierra, Carlos Alberto Barrero, Raul Rodriguez, Yolanda Silva,
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Received 25 March 2003; accepted 4 June 2003



Quantifying *Aotus* monkey cytokines by real-time quantitative RT-PCR

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Expression, polymorphism analysis, reticulocyte binding and serological reactivity of two *Plasmodium vivax* MSP-1 protein recombinant fragments

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Characterizing T-cell receptor γ -variable gene in *Aotus nancymae* owl monkey peripheral blood

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Tissue Antigens.

Tissue Antigens 2003 **62**: 472–482

Hepatitis C virus (HCV) E1 and E2 protein regions that specifically bind to HepG2 cells

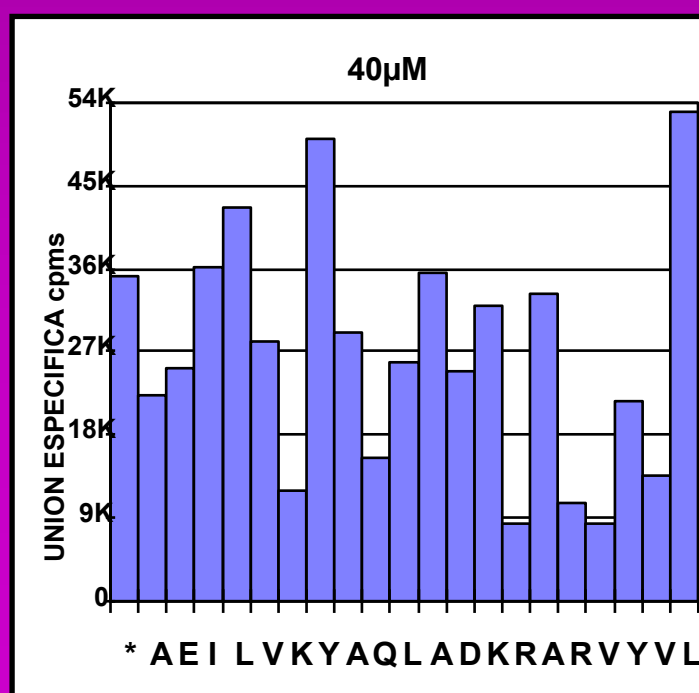
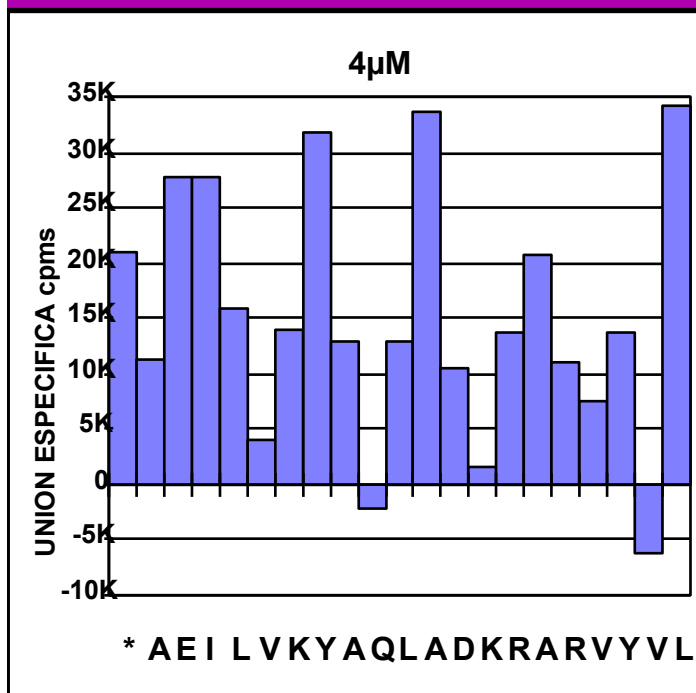
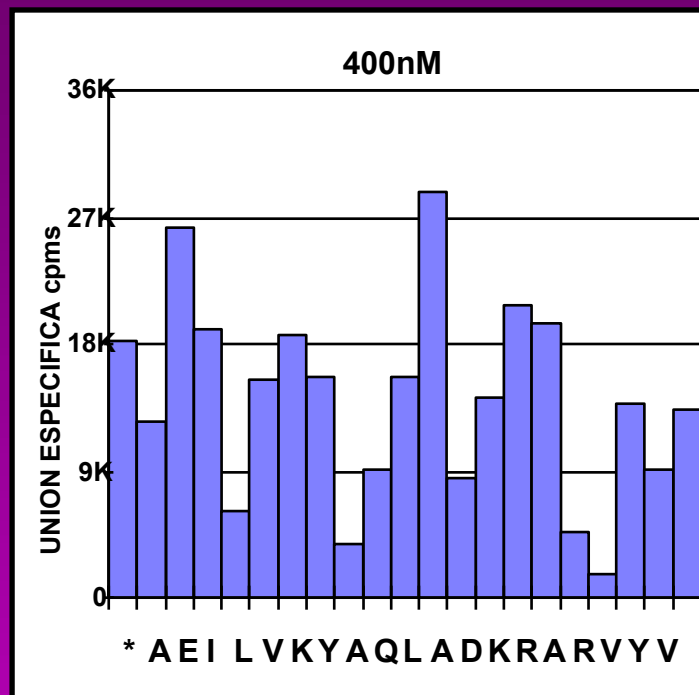
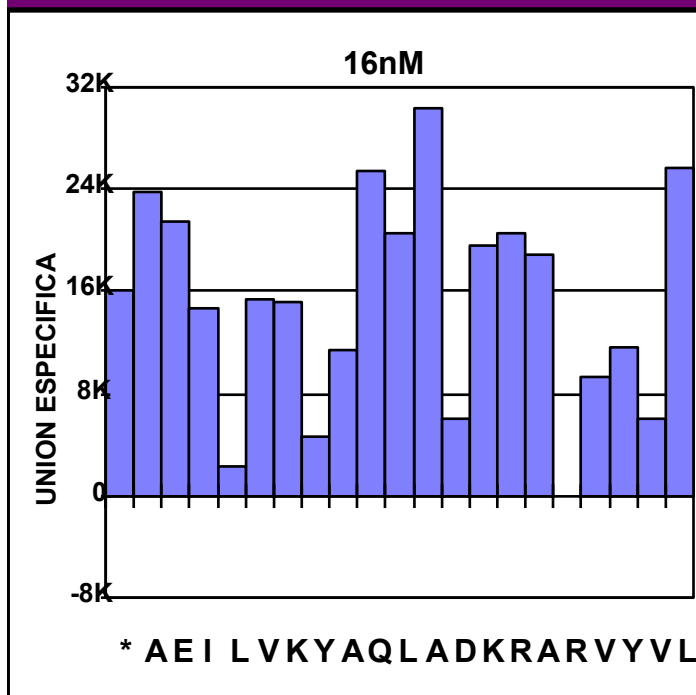
Javier Eduardo García^a, Alvaro Puentes, Jorge Suárez, Ramses López, Ricardo Vera,
Luis Eduardo Rodríguez, Marisol Ocampo, Hernando Curtidor, Fanny Guzmán,
Mauricio Urquiza, Manuel Elkin Patarroyo

PEPTIDE	SEQUENCE	Binding Activity to HepG2 cells 2 %
4931	ETHVTGGSAGHTVSGFVSLLY	
4932	HTVSGFVSLLAPGAKQNVQLY	
4933	APGAKQNVQLINTNGSWHLNY	
4934	INTNGSWHLNSTALNCNDSLY	
4935	STALNCNDSLNTGWLGLFY	
4936	NTGWLGLFYHHKFNSSGCP	
4937	HHKFNSSGCPERLASCRPLTY	
4938	ERLASCRPLTDFDQGWGPISY	
4939	DFDQGWGPISYANGSGPDQR	
4940	YANGSGPDQRPYCWHPYPPKP	
4941	PYCWHPYPPKPCGIVPAKSVC	
4942	CGIVPAKSVCGPVYCFTTSP	
4943	PVYCFTTSPVVVGTTDRSGA	
4944	VVGTTDRSGAPTYSWGEND	
4945	APTYSWGENDDTDFVFLNNTR	
4946	TDVFVFLNNTRPPLGNWFGCTY	
4947	PPLGNWFGCTWMNSTGFTKVY	
4948	WMNSTGFTKVCGAPPCVIGGY	
4949	CGAPPCVIGGAGNNTLHCPTY	
4950	AGNNTLHCPTDCFRKHPDATY	
4951	DCFRKHPDATYSRCGSGPW	
4952	YSRCGSGPWITPRCLVDYPY	
4953	TPRCLVDYPYRLWHYPCTIN	
4954	RLWHYPCTINYTIFKIRMYV	
4955	YTIFKIRMYVGGVEHRLEAA	
4956	GGVEHRLEAACNWTRGERCDY	
4957	CNWTRGERCDLEDNRDRSELSY	
4958	LEDNRDRSELSPLLLTTTQWQY	
4959	PLLLTTTQWQVLPSCFTTLPY	
4960	VLPSCFTTLPALSTGLIHLHY	
4961	ALSTGLIHLHQNIQDVQYLY	
4962	QNIQDVQYLYGVGSSIASWA	
4963	GVGSSIASWAIKWEYVLLF	
4964	ASWAIKWEYVLLFLLLADA	
4965	CONTROL	

PERFIL DE UNION DE PEPTIDOS DERIVADOS DEL CLON MTCY04C12.18c DE *M. tuberculosis*

PEPTIDO	SEQUENCE	1%
11175 ¹	MI ATTRDREGATMI TFRLRLY ²⁰	
11176 ²¹	PCRTI LRVFSRNPLVRGTDRY ⁴⁰	
11177 ⁴¹	LEAVVMLLAVTVSLLTI PFAY ⁶⁰	
11178 ⁶¹	AAAGTAVQDSRSHVYAHQAQ ⁸⁰	
11179 ⁸¹	TRHPATATVI DHEGVI DSNTY ¹⁰⁰	
11180 ¹⁰¹	TATSAPPRTKI TVPARWVNY ¹²⁰	
11181 ¹²¹	GI ERSGEVNAKPGTKSGDRVY ¹⁴⁰	
11182 ¹⁴¹	GI WDSAGQLVDEPAPPARAY ¹⁶⁰	
11183 ¹⁶¹	I ADAAL AAL GLWLSVAAGY ¹⁸⁰	
11184 ¹⁸¹	ALLALTRAI LI RVRNASWQHY ²⁰⁰	
11185 ¹⁹¹	I RVRNASWQHDI DSLFCTQRY ²¹⁰	

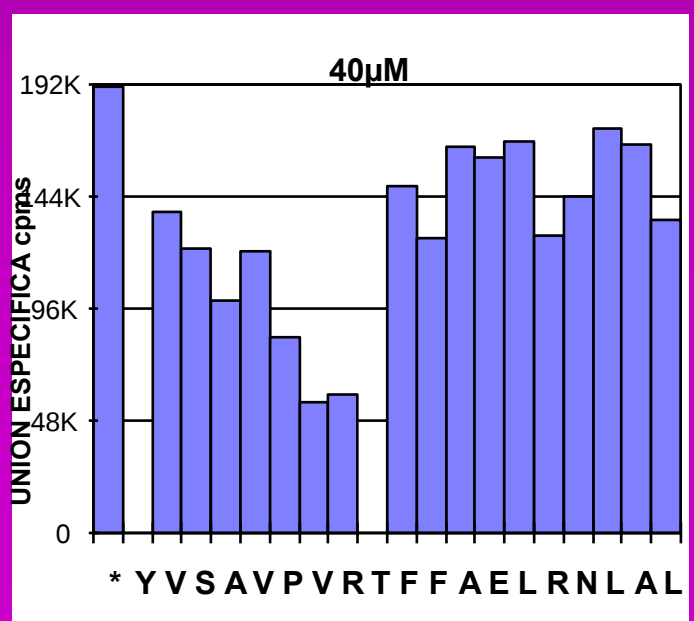
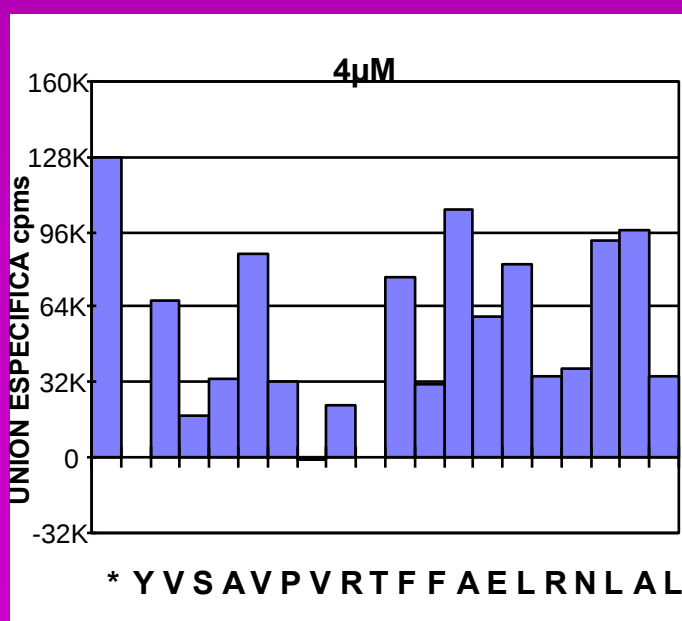
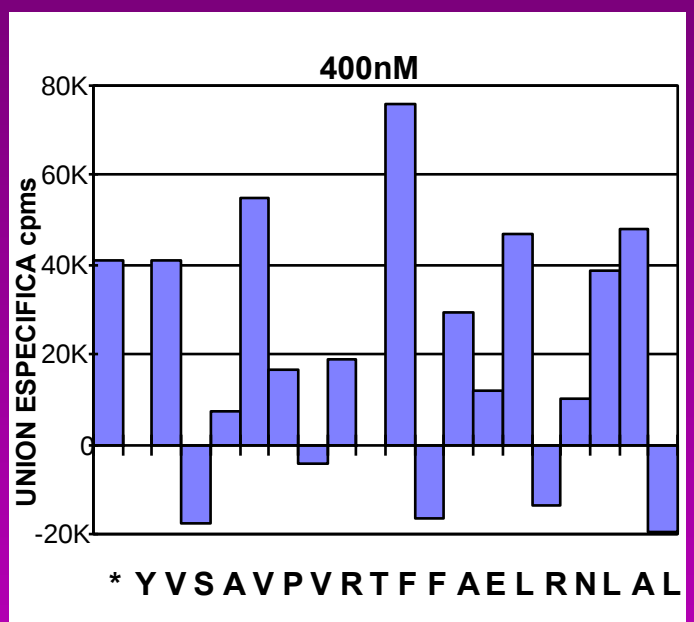
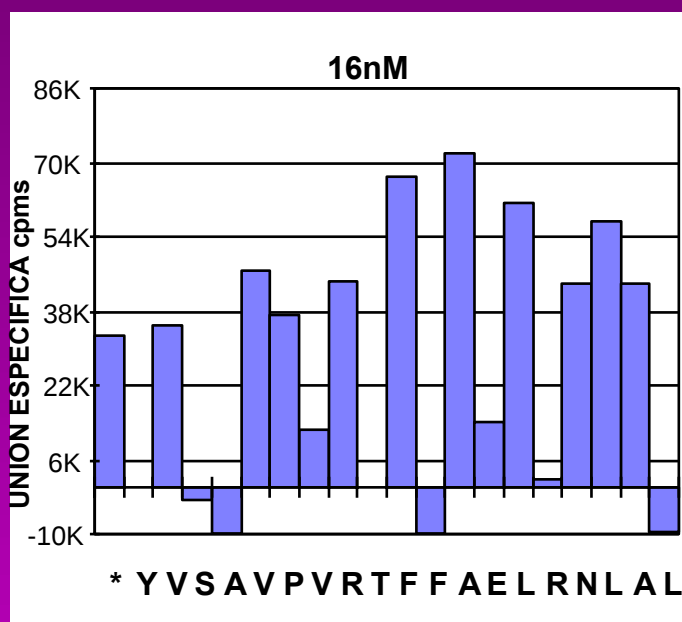
MYCOBACTERIUM TUBERCULOSIS: ANALOGOS DEL PEPTIDO 11060 DE LA PROTEINA PUTATIVA DE MEMBRANA MTCY277.12



PERFIL DE UNION DE PEPTIDOS DERIVADOS DEL CLON MTCY159.20c DE *M. tuberculosis*

PEPTIDO	SEQUENCE		1%
11199	¹ MTNWMLRGLAFAAAMVVLRL	20	
11200	²¹ FQGALINAWQMLSGLISLVL	40	
11201	⁴¹ LLLFAIGGVVWGVMDGRADA	60	
11202	⁶¹ KASPDPPDRRQDLAMTWLLAGY	80	
11203	⁸¹ LVAGALSGAVAWLISLFYKA	100	
11204	¹⁰¹ IYTGGPINELTTFAAFTALI	120	
11205	¹²¹ VFLVGI VGVAVGRWLVDRQLY	140	
11206	¹⁴¹ AKAPVRHHGLAAEHERAADTY	160	
11207	¹⁶¹ DVFS AVRADDSP TGEMQVAQY	180	
11208	¹⁸¹ PEAQTA AVATVEREAPTEVI	200	
11209	²⁰¹ RTTESDTPTEVIRTDTEADQY	220	
11210	²¹¹ VI RTDTEADQTKPGDEPKKDY	230	

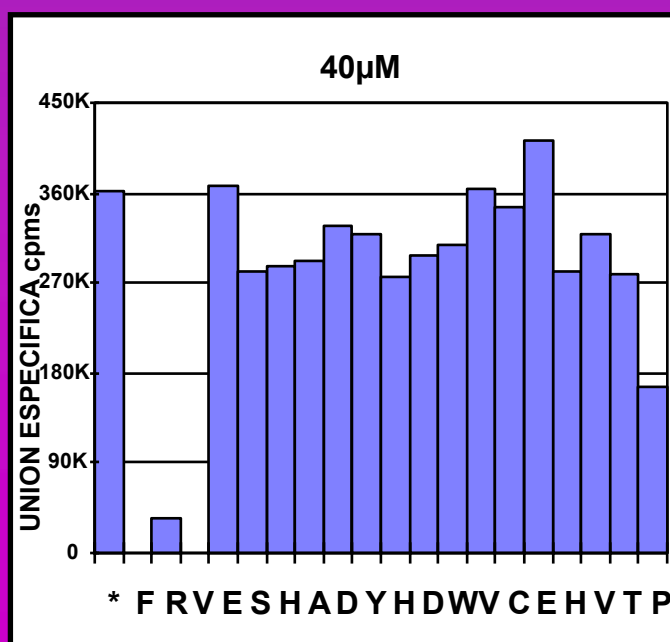
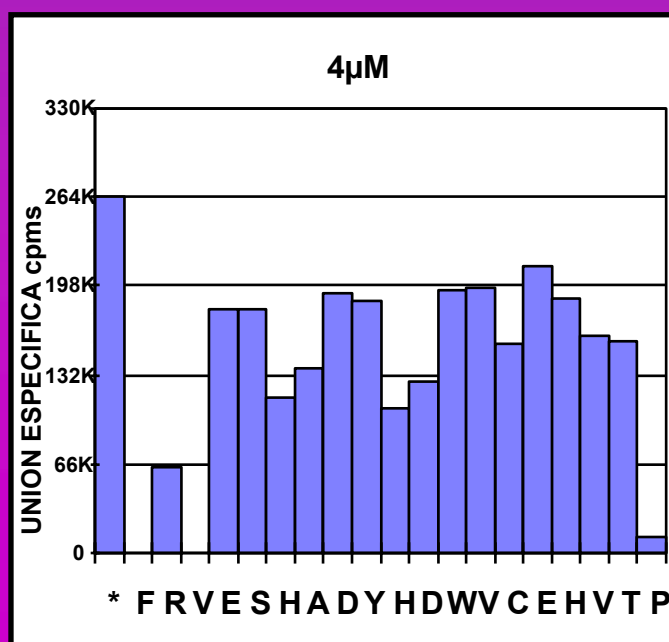
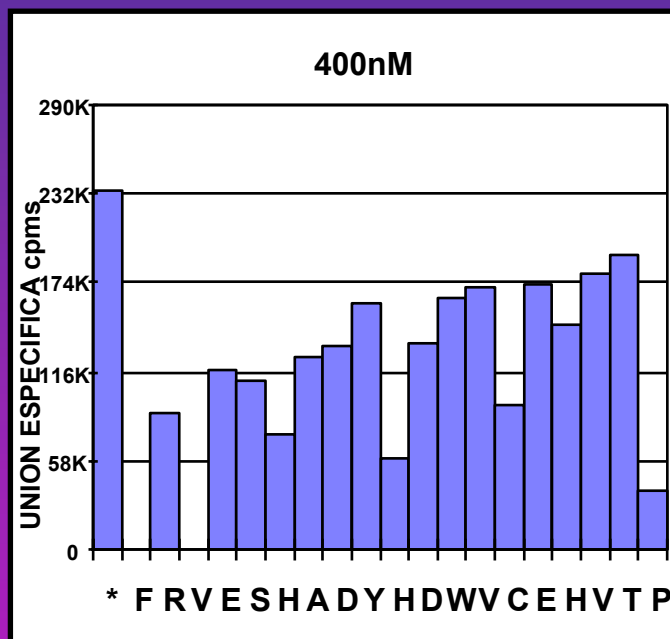
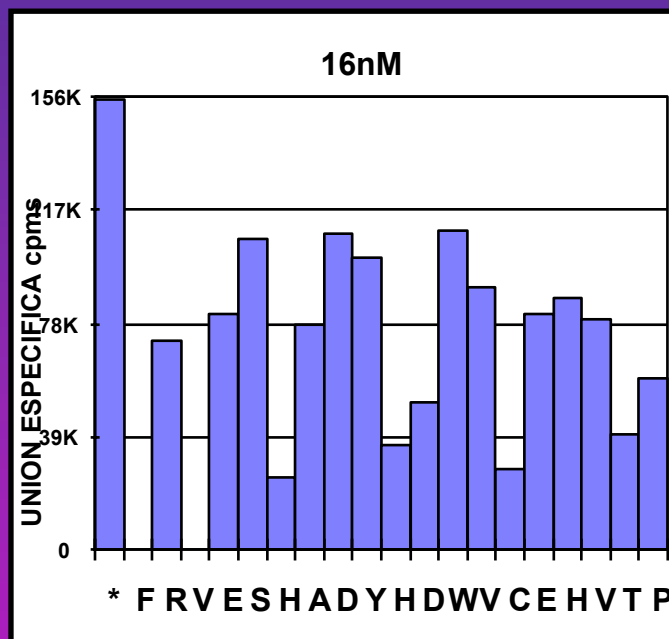
MYCOBACTERIUM TUBERCULOSIS: ANALOGOS DEL PEPTIDO 11076 DE LA PROTEINA PUTATIVA DE MEMBRANA MTCY277.12



PERFIL DE UNION DE PEPTIDOS DERIVADOS DEL CLON MTCY277.12 DE *M. tuberculosis*

PEPTIDO	SEQUENCE		1%
11059	¹ MSQCF AVKGI GGADQATLGSY	20	
11060	²¹ AEI LVKYAQLADKRARVYVL	40	
11061	⁴¹ VST WL VVWGI WHVYFVEAVF	60	
11062	⁶¹ PNAI L WL HYAASYEF GFVR	80	
11063	⁸¹ RGL GGELI RMLTGDHFFAGAY	101	
11064	¹⁰¹ YTVLWTSI TVWLI ALAVVW	120	
11065	¹²¹ LI LSTGNRSERRI MLALLVPY	140	
11066	¹⁴¹ VLPFAFSYAI YNPHPELFGM	160	
11067	¹⁶¹ TALVAFSI FLTRAHTSRTRVY	180	
11068	¹⁸¹ I LSTLYGLTMAVLALI HEAI	200	
11069	²⁰¹ PLEFALGAVLAI I VLSKNATY	220	
11070	²²¹ GATRRI CTALAI GPGTVSVLY	240	
11071	²⁴¹ LLAVVGRRDI ADQLCAHI PHY	260	
11072	²⁶¹ GMVENPWAVATTPQRVLDYI	280	
11073	²⁸¹ FGRVESHADYHDWVCEHVTP	300	
11074	³⁰¹ WFNLDWI TSAKLVAVVGFRAY	320	
11075	³²¹ LFGAFL LGLLFFVATTSMI RY	341	
11076	³⁴¹ YVSAVPVRTFFAELRGNLAL	360	
11077	³⁶¹ PVLASALLVPLFI TAVDWTRY	380	
11078	³⁸¹ WWVMI TLDVAI VYI LYAI DR	400	
11079	⁴⁰¹ PEI EQPPSRRNVQVFVCVVLY	420	
11080	⁴¹⁶ VCVVLVLAVI PTGSANNI GRY	435	

MYCOBACTERIUM TUBERCULOSIS: ANALOGOS DEL PEPTIDO 11073 DE LA PROTEINA PUTATIVA DE MEMBRANA MTCY277.12

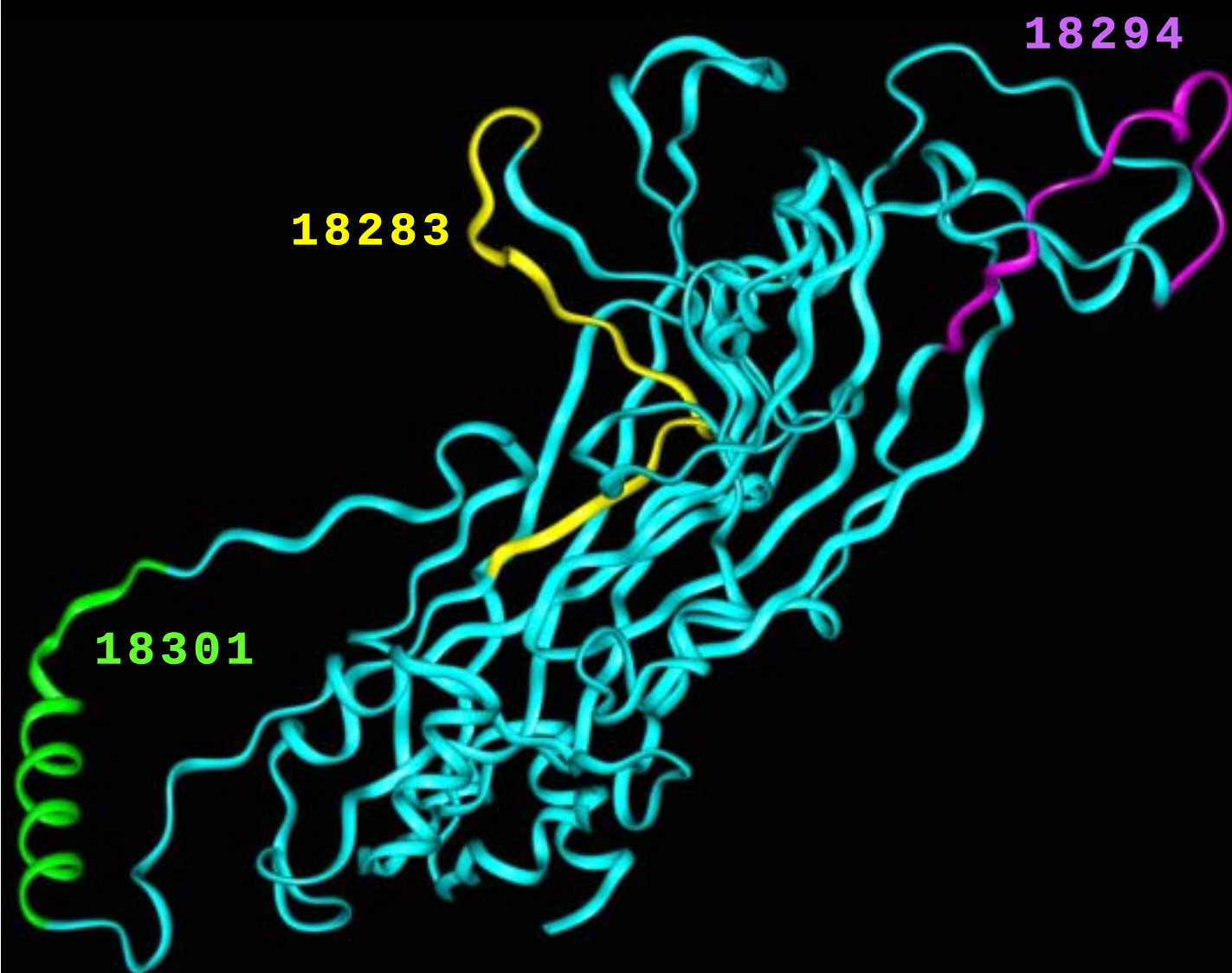


HUMAN PAPILLOMAVIRUS TYPE 16 AND 18 L1 PROTEIN PEPTIDE BINDING TO VERO AND HELA CELLS INHIBITS THEIR VLPS BINDING

Ricardo VERA-BRAVO, Marisol OCAMPO, Mauricio URQUIZA, Javier E. GARCÍA, Luis E. RODRÍGUEZ, Alvaro PUENTES, Ramses LÓPEZ, Hernando CURTIDOR, Jorge E. SUÁREZ, Elizabeth TORRES, Fanny GUZMÁN, Diana DÍAZ, Jimena CORTES, María M. BRAVO, Alba L. CÓMBITA, Oscar OROZCO and Manuel E. PATARROYO*

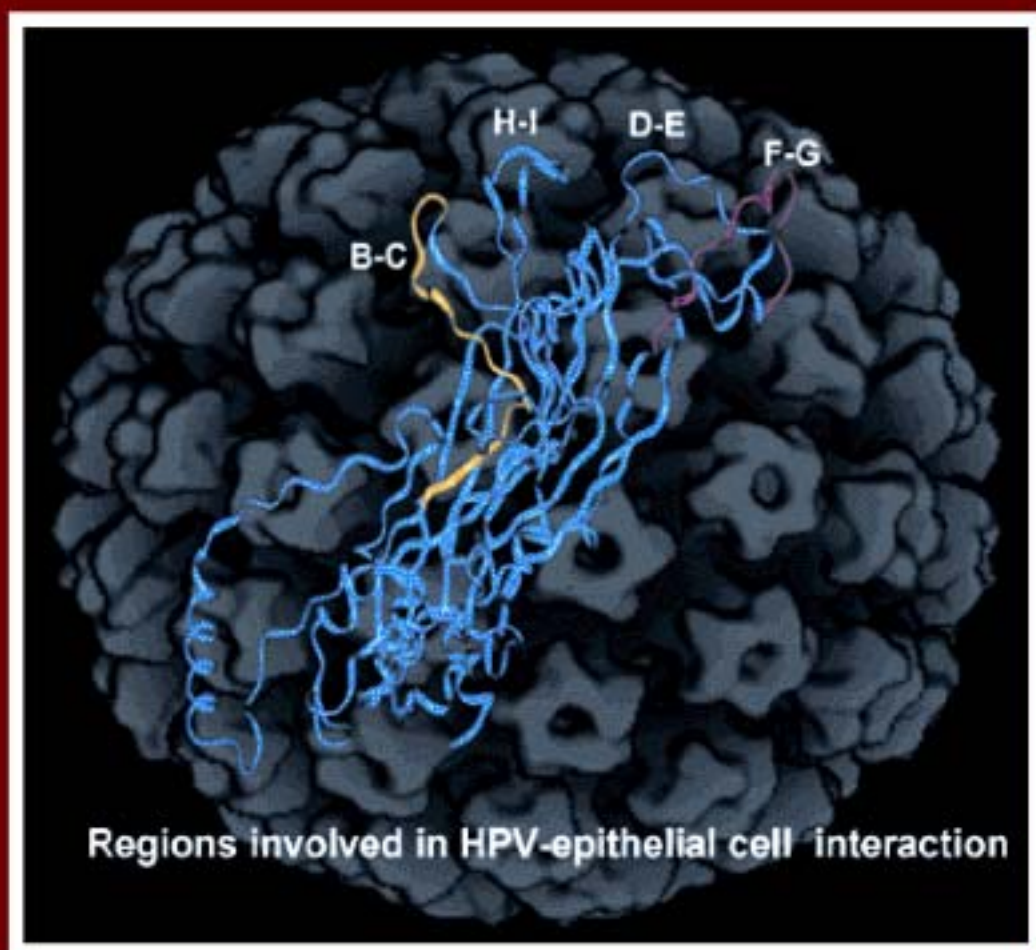
Fundación Instituto de Inmunología de Colombia, Instituto Nacional de Cancerología and Universidad Nacional de Colombia, Bogotá, Colombia

PEPTIDE NUMBER	SEQUENCE	High Specific Binding Activity (VERO CELLS)	High Specific Binding Activity (HeLa CELLS)
18306	1 MCLYTRVLILHYHLLPLYGP 20		
18307	21 LYHPRPLPLHSILVYMHII 40		
18308	41 ICGHYILFLRNVNVEPIFLI 60		
18309	61 QMALWRPSDNTVYLPPPSVA 80		
18310	81 RVVNTDDYVTPTSIFYHAGS 100		
18311	101 SRLTVGNPYFRVPAGGGNK 120		
18312	121 QDIPKVSAYQYRVFRVQLPD 140		
18313	141 PNKFGLPDTSIYNPETQRLV 160		
18314	161 WACAGVEIGRGQPLGVGLSGY 180		
18315	181 HPFYNKLDDESSHAATSNV 200		
18316	201 SEDVRDNVSDYKQTQLCIL 220		
18317	221 GCAPAIGEHWAKGTACKSRPY 240		
18318	241 LSQGDCCPLELKNTVLEDGDY 260		
18319	261 MVDGTGYGAMDFSTLQDTKCE 280		
18320	281 VPLDICQSICKYPDYLOMSA 300		
18321	301 DPYGDSMFFCLRREQLFARH 320		
18322	321 FWNRACTMGDTVPQSLYIKG 340		
18323	341 TGMSPASPGSCVYSPSPSGSI 360		
18324	361 VTSDSQLFNKPYWLHKAQGH 380		
18325	381 NNGVCWHNQLFVTVDTPSY 400		
18326	401 TNLTICASTQSPVPGQYDAT 420		
18327	421 KFKQYSRHHVEEYDLQFIFQL 440		
18328	441 CTITLTADVMSYIHSNMSSI 460		
18329	461 LEDWNFGVPPPTTSLVDY 480		
18330	481 RFVQSVAITCQKDAAPAENKY 500		
18331	501 DPYDKLKFVNVDLKEKFSLD 520		
18332	521 LDQYPLGRKFLVQAGLRKPK 540		
18333	541 TIGPRKRSAPSATTSSKPAKY 560		
18334	561 APSATTSSKPAKRVRVRARKY 580		



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Herpes Virus Saimiri

Peptide Sequence and Binding Activity of Probable Antigen 75

Curtidor H. et al. Manuscript in preparation

PEPTIDE NUMBER	SEQUENCE	Specific Binding Activity (VERO) 2%	Specific Binding Activity (OMK) 2%
11720	1 MAANRHGGHLLPLEGLAAPT Y 20		
11721	21 HRTVIYYAKCDFSPSEERC V 40		
11722	41 NRFTGRPGPLTLMRDRSNVE Y 60		
11723	61 HMLVITVKTEDEDRNTAEQNY 80		
11724	81 QARARSRSQETKKILTYLLAP Y 100		
11725	101 QIDYNPLTLDSLHDHNLGQRA 120		
11726	121 VIFSYPNLHQRLTTSALEL 140		
11727	141 QRACKTLKLHLSILRIESGRHY 160		
11728	161 FTSKAVQYVVVENEDDILTAA 180		
11729	181 LIGENNLYQINTETLSATRL 200		
11730	201 VTDALSWAEYKPLDVTCTNAV 220		
11731	221 MKPPREGVQPMCLVASSPVSY 240		
11732	241 DYNFNKYLMITPSCTAISLH 260		
11733	261 MGHPYPRVSQGIHVVHSSTR 280		
11734	281 TMGRQCCDYLFHSSLLPNGN 300		
11735	301 TTAFAAGYVVTSTTGSGSIP 320		
11736	321 TSVNEQILHAASAQGRSLNNY 340		
11737	341 LGVPVVSQGLKPLPRCSEVPY 360		
11738	361 NNVITHVSQRLRTASERDLNLY 380		
11739	381 CRVRAGQFVAAVGAYDPTSG 400		
11740	401 PDKSPYLYRDSIDSLNRAIQ 420		
11741	421 ATKLFYQMCETPCVSSYQRE 440		
11742	441 FGSCSTLHHLLALVSPKGMTY 460		
11743	461 VHISRLPEEITKALRSVPVLY 480		
11744	481 EEDTVCAFVSQYFFNCFSSQ 500		
11745	501 LFLVIDDKVKITTPSGQIHFTY 520		
11746	521 DILKKAGNLCGAPVYILGRT 540		
11747	541 CNDIGIHCVNDLYHPRDLSV 560		
11748	561 LDSQATSSMTLTVQPHASVVY 580		
11749	581 SATLQPQEPHEEDESIDWAMY 600		
11750	601 FGTSSITISNLSHPAVASKSY 620		
11751	621 NIIRRLDRCGNGLIAQQPGIY 640		
11752	641 GPSDAPVSDYAIICDSSMFP 660		
11753	661 ARLENDASIKKISKQEAQRY 680		
11754	681 AYAQIHKWFGTEKLFINTIS 700		
11755	701 AKVIALGEQAYKLSRNPIVG 720		
11756	721 VKYAI AEAITNIMFGPDCVL 740		
11757	741 EDITLTVAAHWNKQETAALY 760		
11758	761 RVLFACKEMCRELNVNLSITY 780		
11759	781 DASDSRDTPIQDTDAANTVY 800		
11760	801 VTASARVTSIERITPALKKAY 820		
11761	821 ENALVHVCLSKELTLGSGSVFY 840		
11762	841 ENSFTAFFSSHLPDLDTSKLRY 860		
11763	861 DMFYAVKHLISKNLVVAGHD 880		
11764	881 ISDGGILTTAAEMCFASTFGY 900		
11765	901 VTVNLPSALPALMYLVSETP 920		
11766	921 GALLEVPKEHLSTVTTLLEY 940		
11767	941 RGLTWYAVGTVNNVKNLSIY 960		
11768	961 DNGTHLLTESINILNSKWMSY 980		
11769	981 YAEESFETCEPHIESMYRND 1000		
11770	1001 YGNAMDLKHLEDLCTHKPL 1020		
11771	1021 QLYTCPSHPVSVAVLTFPGC 1040		
11772	1041 PDPVATLQAFANVGFLSYPI 1060		
11773	1061 STEFLLQGNNLNAFSLAVSY 1080		
11774	1081 GSSAFEEEGTGTRIAIYTLL 1100		
11775	1101 QCDLAKNCLKEFFQRPDTLSY 1120		
11776	1121 LCCGELGTQLLAACQVVGDTY 1140		
11777	1141 HPSRGDISSNPESWTLELEPY 1160		
11778	1161 NASKHYESLWLNFBHVPQTIK 1180		
11779	1181 SIILQALRGITFQDGLWQVLY 1200		
11780	1201 GLRYKHDAQEYIMQQNGTIA 1220		
11781	1221 MSYHSAKINPYLYAMHYPRN 1240		
11782	1241 PSGNSSVAGICSKNGRHLALY 1260		
11783	1261 LVEPALSFHTWQWQHIPKPLY 1280		
11784	1281 VTSPWALMYQCMFLWCVKE 1296		

Constructing a useful tool for characterizing amino acid conformers by means of quantum chemical and graph theory indices

Constanza Cárdenas^{a,b,*}, Mateo Obregón^a, Eugenio-José Llanos^{a,b},
Eduardo Machado^a, Hugo-Javier Bohórquez^a, Jose-Luis Villaveces^b,
Manuel-Elkin Patarroyo^{a,b}

21 índices de **mecánica cuántica**

19 índices **grafo-teóricos**

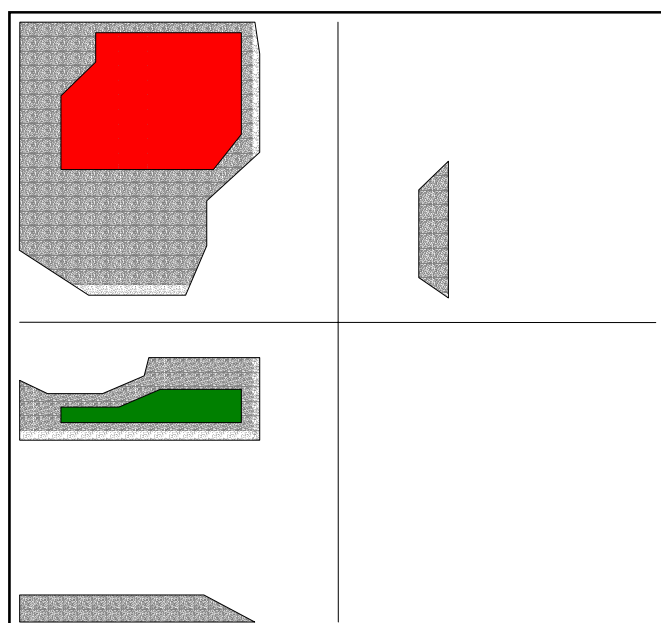
Ala-X-Ala

NH₂-X-CHO

• 2 conformaciones (alfa y beta)

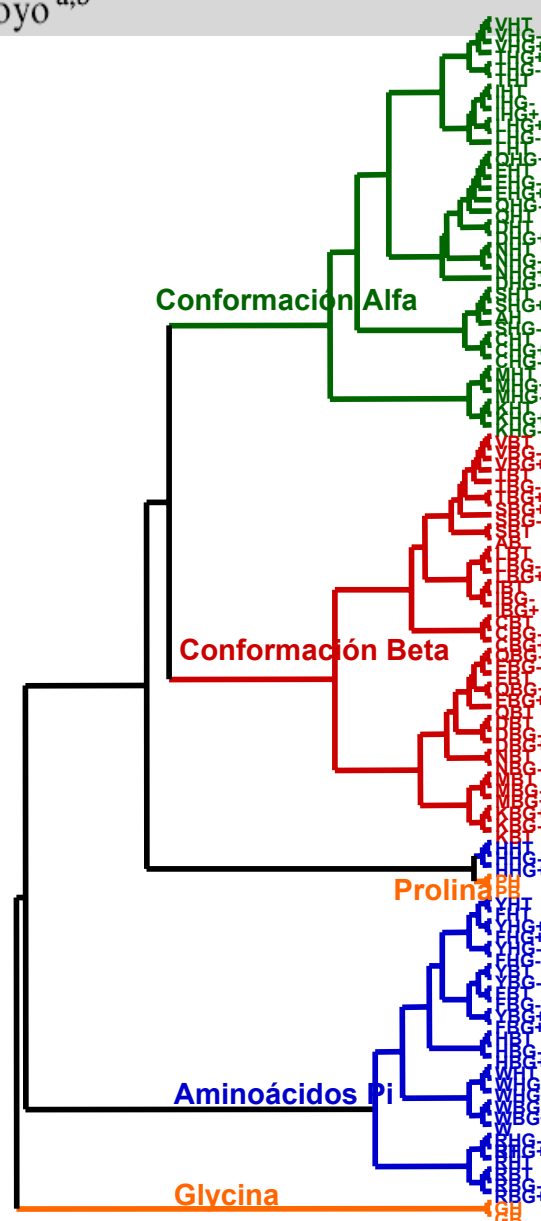
• 3 rotámeros (g⁺, g⁻, t)

540 moléculas



φ

ψ



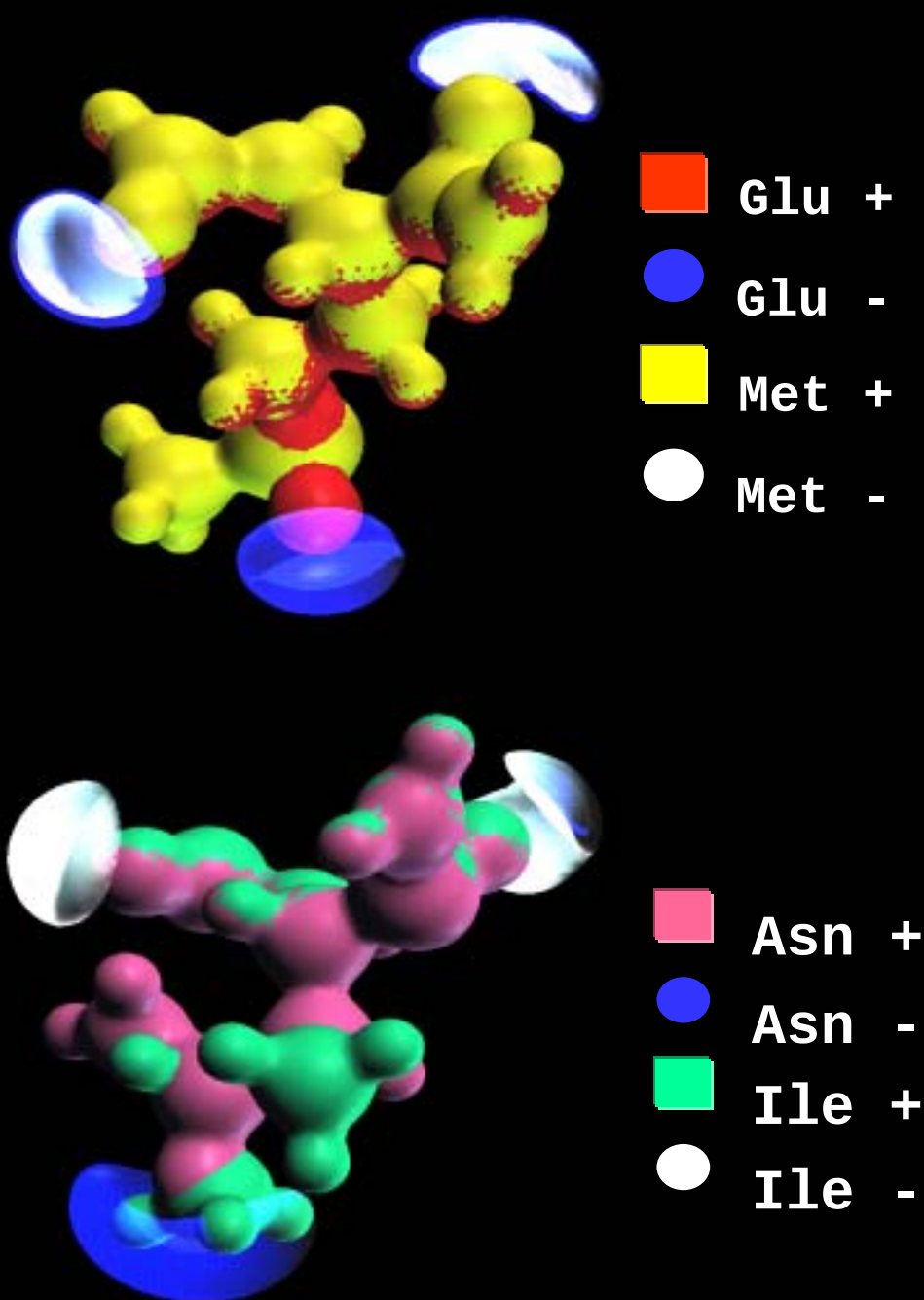
Electronic Energy and Multipolar Moments Characterize Amino Acid Side Chains into Chemically Related Groups

Hugo J. Bohórquez,* Mateo Obregón, Constanza Cárdenas, Eugenio Llanos, Carlos Suárez, José Luis Villaveces, and Manuel Elkin Patarroyo

Fundación Instituto de Immunología de Colombia-FIDIC, Carrera 50 No. 26-00, Bogotá, Colombia

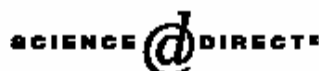
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Quantum chemical analysis explains hemagglutinin peptide–MHC Class II molecule HLA-DR β 1*0101 interactions

Constanza Cárdenas^{a,b,1}, José Luis Villaveces^{b,1}, Hugo Bohórquez^a, Eugenio Llanos^b, Carlos Suárez^a, Mateo Obregón^a, Manuel Elkin Patarroyo^{a,*}

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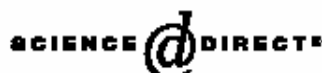
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A comparative study of MHC Class-II HLA-DR β 1*0401-Col II and HLA-DR β 1*0101-HA complexes: a theoretical point of view

Constanza Cárdenas^{a,b,*,1}, José Luis Villaveces^b, Carlos Suárez^{a,1}, Mateo Obregón^{a,1}, Marysol Ortiz^{a,1}, Manuel Elkin Patarroyo^{a,1}

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Synthetic vaccines - Emerging principles

1. Plasmodia (falciparum & vivax) use CONSERVED as well as VARIABLE sequences to bind to host cells.
2. CONSERVED binding sequences are NOT antigenic, but VARIABLE sequences are
3. CONSERVED binding sequences are NOT immuno-genic in ANY of the tested species when administered as monomers, polymerized, associated with T helper epitopes or presented as MAPs. While VARIABLE sequences can be immunogenic they are strain-specific
4. To overcome this problem the critical contact residues in the binding interaction have to be identified and changed
5. The critical contact residues have to be changed for aminoacids that ARE NOT commonly used during the mutational process in the variable sequences
6. Physicochemical characteristics such as size, charge and volume have to be considered. Therefore $A \leftrightarrow S$; $P \leftrightarrow D$; $F \leftrightarrow R$; $H \leftrightarrow L$, $K \leftrightarrow M$; $N \leftrightarrow I$; $Q \leftrightarrow E$; $C \leftrightarrow T$; $Y \leftrightarrow W$; can be exchanged. G and C have special properties.
7. There must be a $23.5 \text{ \AA} + 3.5 \text{ \AA}$ distance between those residues fitting into Pockets 1-9 of Class II molecules.
8. All the changes made to the CONSERVED binding peptides are designed to achieve a perfect fit with the MHC molecules and the TCR
9. Structural considerations together with affinity, biological and Immunological activities should guide the design of synthetic peptide vaccines.
10. This methodology elicits protective immunity against NON ANTIGENIC, NON IMMUNOGENIC CONSERVED BINDING PEPTIDES, providing a new and powerful tool for the rational development of synthetic vaccines

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NMR y Estructura 3D



**Síntesis
Química de
Macro-
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