

# **SIMIAN MALARIA IN MALAYSIA WITH SPECIAL REFERENCE TO *PLASMODIUM KNOWLESI***

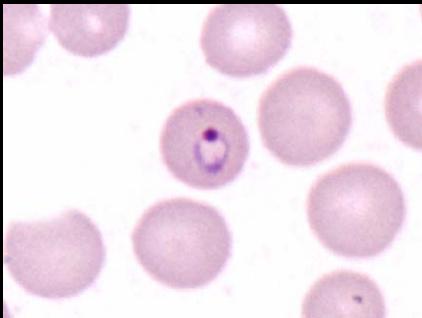


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# INTRODUCTION

- 1<sup>st</sup> natural infection of *P. knowlesi* in humans was reported from Pahang - 1965
- 2<sup>nd</sup> case was reported from Johore in 1971.

# *Plasmodium knowlesi*



*Macaca fascicularis*, *M. nemestrina*, *Pryesbytis melalophos*

# Vectors – Peninsular Malaysia

## *Anopheles hackeri*

- natural vector of *P. knowlesi*
- highly zoophagic

## *Anopheles cracens* (as *An. balabacensis*)

- vector of *P. inui* and *P. cynomolgi*

## *Anopheles latens* (as *An. leucosphyrus*)

- vector of *P. inui*

REF: Wharton *et al* 1961, 1962; Cheong *et al* 1965

# INTRODUCTION

- In 2004 Singh *et al* reported a large focus of *P. knowlesi* in Sarawak Malaysian Borneo  
(Lancet 2004)



Is *P. knowlesi*  
occurring only  
in Malaysian  
Borneo ?



# **GENERAL OBJECTIVES**

- To study simian malaria in humans and non human primates
- To elucidate the vectors of simian malaria

# MATERIALS & METHODS

- Samples received since July 2005
- Extraction of DNA from whole blood or blood film
- Nested PCR following protocol of Singh et al 2004

# Species specific primers used in the Nested-PCR Assay

rPLU1&5

Human malaria  
*P. falciparum*  
*P. vivax*  
*P. malariae*

Simian malaria  
*P. knowlesi*

# CASES BY MICROSCOPY AND PCR JULY 2005-DEC. 2007

	Cases detected by microscopy					Cases detected by PCR
PCR results	Pf	Pv	Pm	Pf+Pm	Pm+Pv	
Pf	4	1	1			6
Pv		6	3	1	1	11
Pm			6			6
Pk	2	1	58			61
Pf+Pm			1			1
Pf+Pk			1			1
Pv+Pk		1	1			2
Pk+Pm			2			2
Total	6	9	73	1	1	90

2008

Jan- Aug.

**MALARIA CASES BY MICROSCOPY**

<b>PCR</b>	<b>Pf</b>	<b>Pv</b>	<b>Pm</b>	<b>Pf+Pv</b>	<b>Pm+Pv</b>	<b>Pf+Pm</b>	<b>Negative</b>	<b>Cases by PCR</b>
<b>Pf</b>	<b>2</b>		<b>2</b>			<b>1</b>		<b>5</b>
<b>Pv</b>		<b>4</b>	<b>12</b>		<b>1</b>		<b>5</b>	<b>22</b>
<b>Pm</b>			<b>2</b>					<b>2</b>
<b>Pk</b>	<b>3</b>		<b>40</b>	<b>2</b>		<b>1</b>		<b>46</b>
<b>Pf+Pv</b>				<b>1</b>				<b>1</b>
<b>Pf+Pk</b>	<b>1</b>		<b>2</b>	<b>1</b>		<b>1</b>		<b>5</b>
<b>Pv+Pk</b>			<b>12</b>	<b>1</b>	<b>1</b>		<b>1</b>	<b>15</b>
<b>Pm+Pk</b>			<b>1</b>					<b>1</b>
<b>Pv+Pf+Pk</b>							<b>1</b>	<b>1</b>
<b>Pf+Pm+Pk</b>						<b>1</b>		<b>1</b>
<b>Total</b>	<b>6</b>	<b>4</b>	<b>71</b>	<b>5</b>	<b>2</b>	<b>4</b>	<b>7</b>	<b>99</b>

# CASES OF *P. KNOWLESI* IN P. MALAYSIA



# NON HUMAN PRIMATES





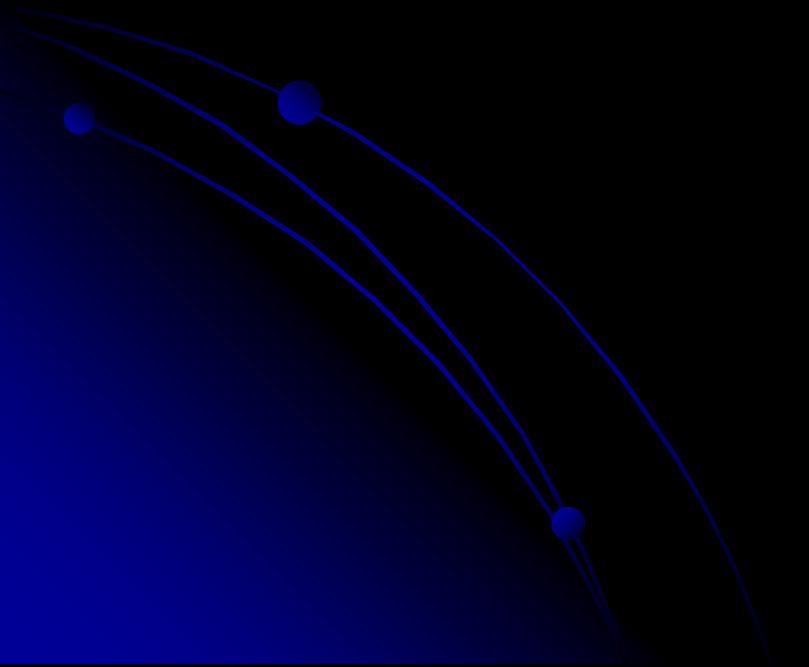
16/08/2007

# Species specific primers used in the Nested-PCR Assay

rPLU1&5

Simian malaria

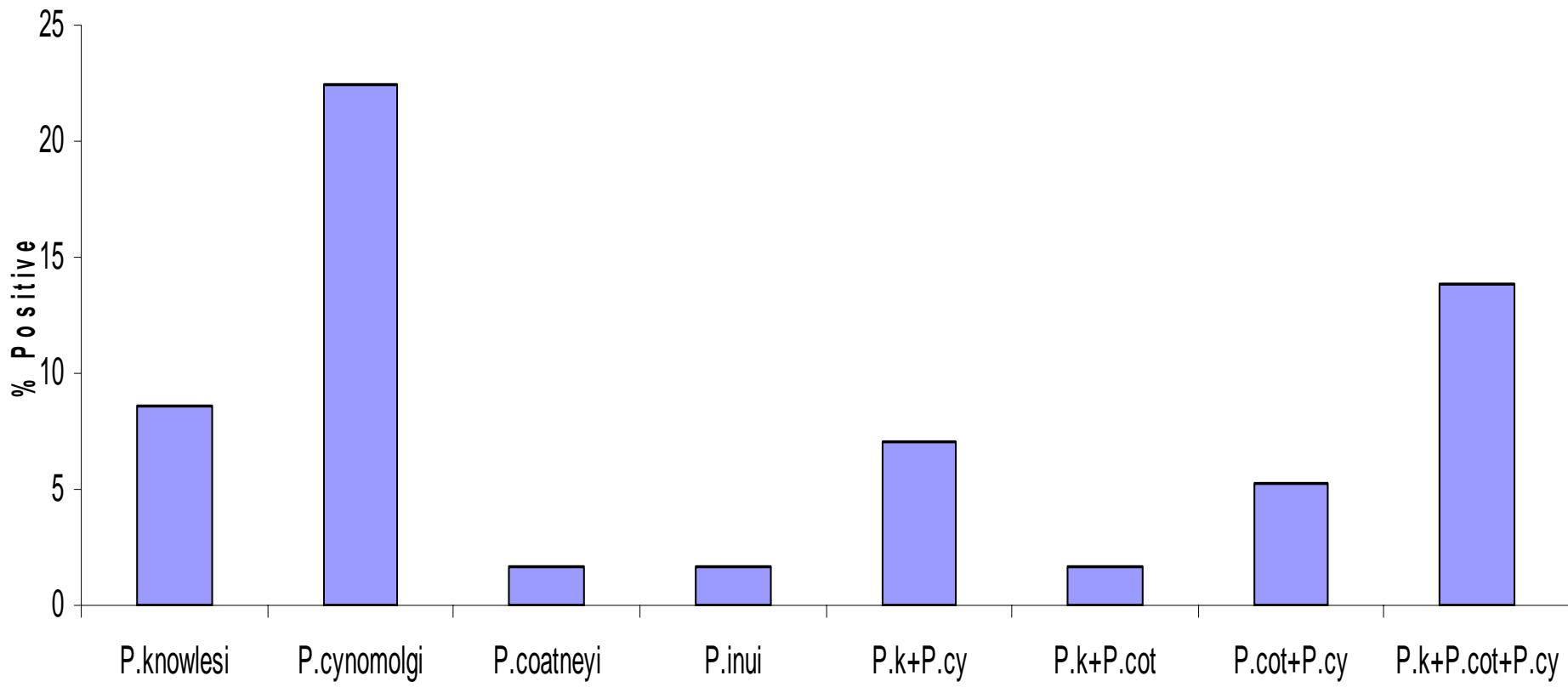
*P. knowlesi*  
*P. coatneyi*  
*P. inui*  
*P. fieldi*  
*P. cynomolgi*  
*P. fragile*



# RESULTS OF MONKEY SAMPLES

Locality	Samples collected	BFMP Positive (%)	PCR Positive Genus (%)
Selangor	60	0	0 (0)
Kuala Lipis	104	102 (98.1)	95 (91.3)
Temerloh	11	11 (100)	11 (100)
Kuala Lumpur	29	2 (6.9)	2 (6.9)
Total	214	115 (56.7)	108 (53.2)

# Malaria Species in Monkeys by PCR



# MOSQUITO COLLECTION STUDY SITE





05/09/2007









15/08/2007

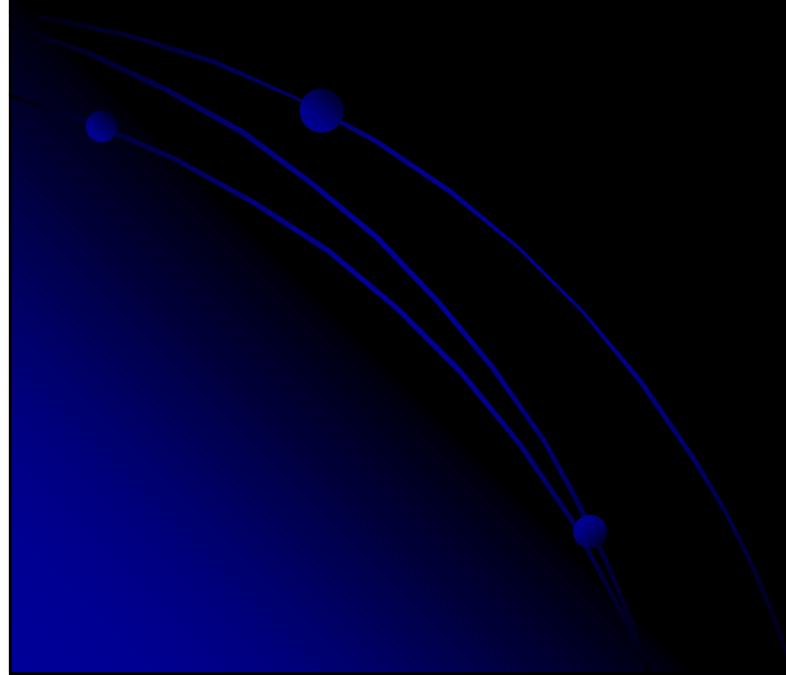
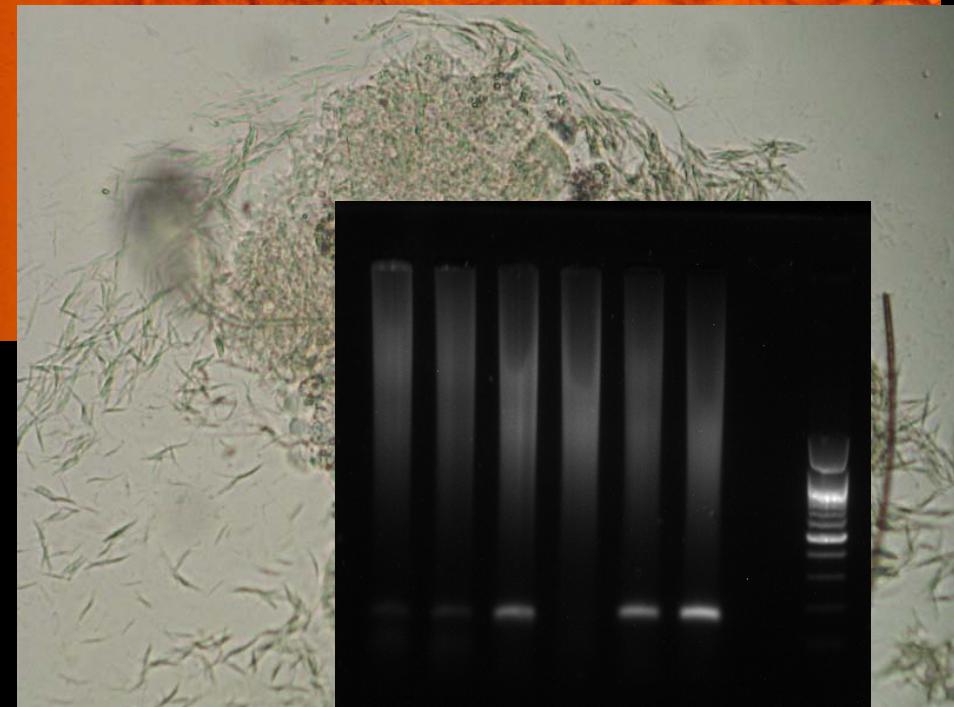


# RESULTS

## Predominant *Anopheles* species in study site Aug 2007-Aug 2008

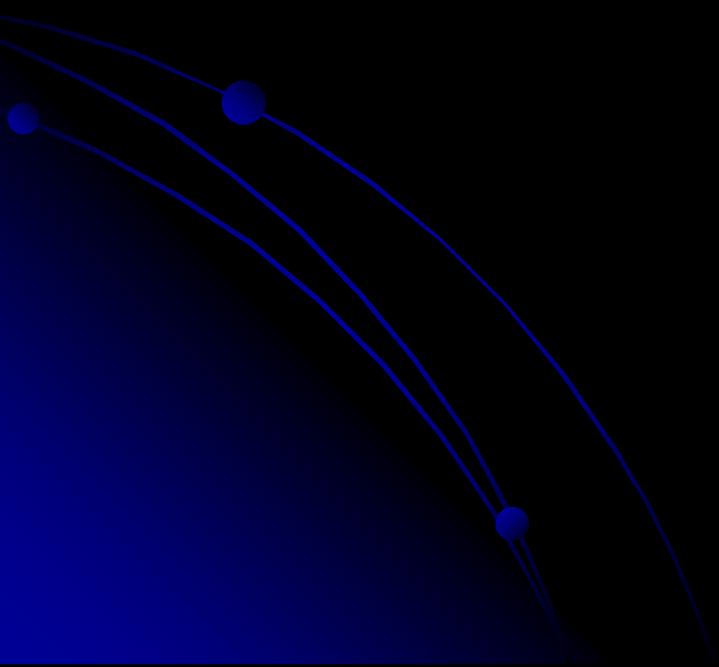
Anopheles species	Mela		Sg Ular	Total (%)
	BLC	MBT		
<i>An.aconitus</i>	0	0	6	6 (0.46)
<i>An. barbirostris</i> gr	10	11	4	25 (1.93)
<b><i>An. cracens</i></b>	<b>186</b>	<b>77</b>	<b>584</b>	<b>847 (65.55)</b>
<i>An. hyrcanus</i> gr	34	15	1	50(3.86)
<b><i>An. kochi</i></b>	<b>2</b>	<b>51</b>	<b>1</b>	<b>54 (4.17)</b>
<b><i>An. maculatus</i></b>	<b>99</b>	<b>1</b>	<b>163</b>	<b>263 (20.35)</b>
<i>An. philippinensis</i>	6	0	6	12 (0.92)
<i>An. pujutensis</i>	0	1	0	1 (0.07)
<i>An. separatus</i>	2	1	5	8 (0.61)
<i>An. tesselatus</i>	7	10	5	22 (1.70)
<i>An. umbrosus</i>	1	1	0	2 (0.15)
<i>An. vagus</i>	1	1	0	2 (0.15)
<b>Total</b>	<b>348</b>	<b>169</b>	<b>775</b>	<b>1292</b>

# RESULTS

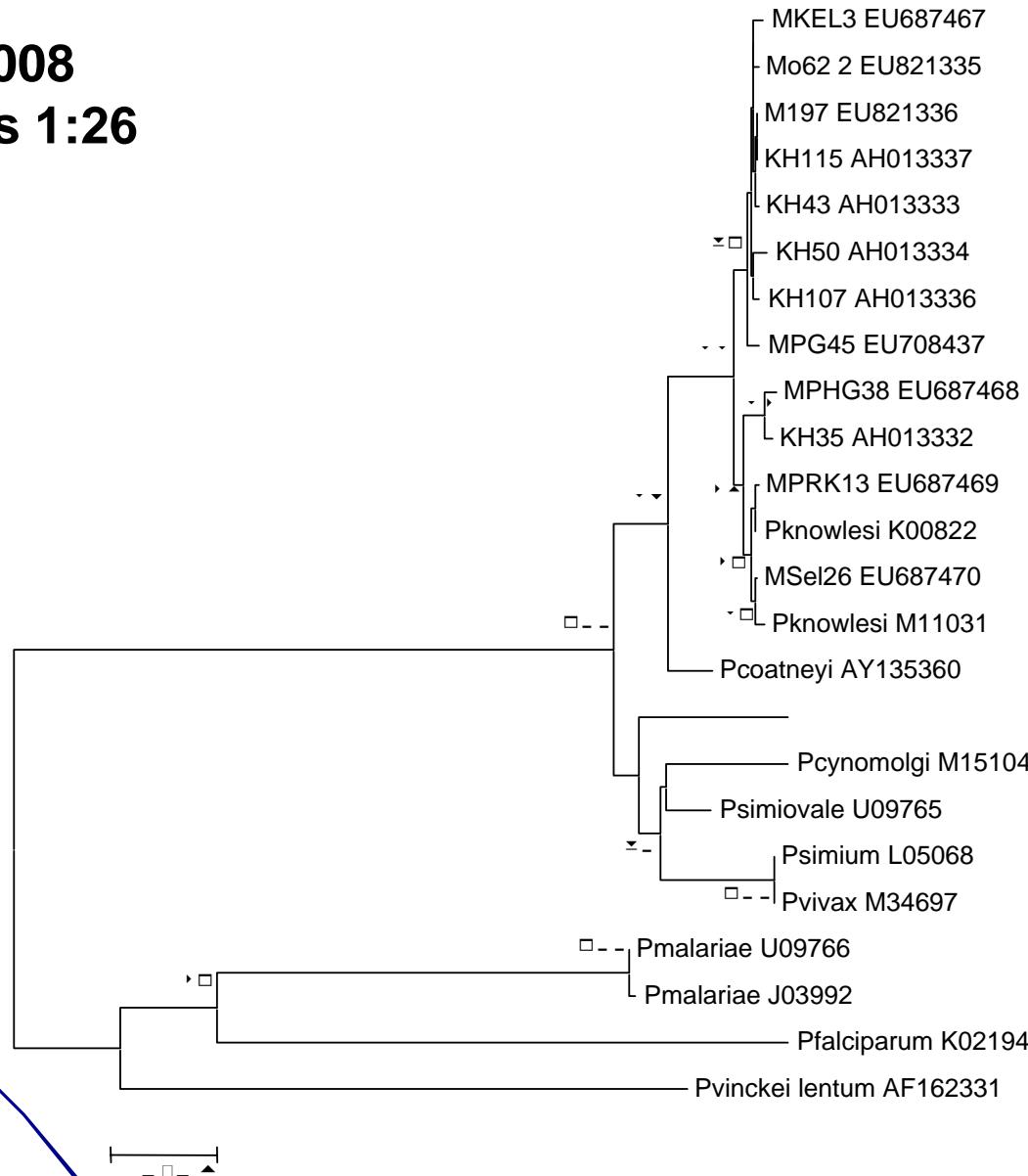




To verify species of *Plasmodium* found,  
cloning and sequencing of the  
circumsporozoite protein (*csp*) genes  
were conducted...

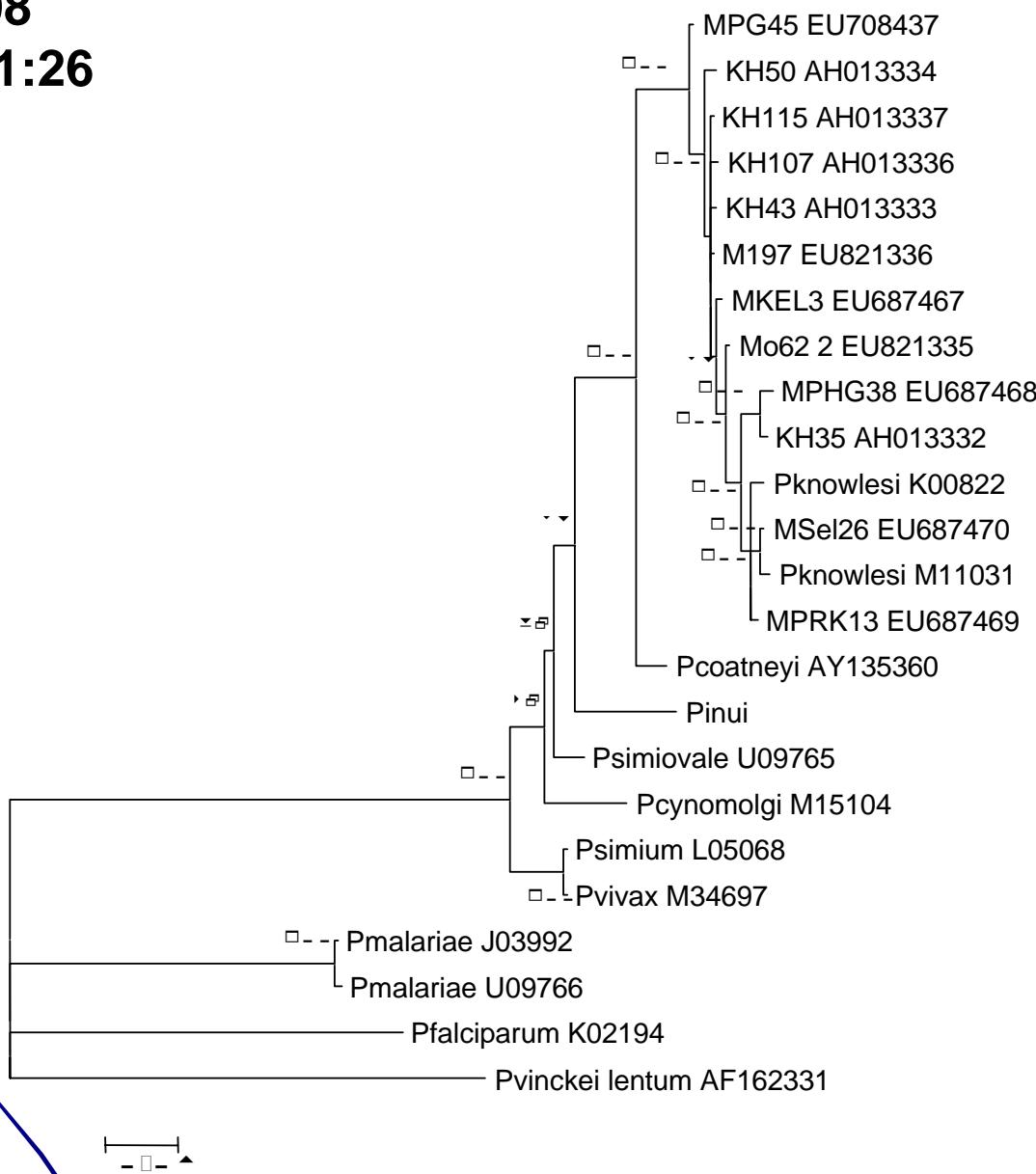


Vythilingam et al 2008  
Parasites & Vectors 1:26



Phylogenetic tree based on the non-repeat region of the circumsporozoite (csp) genes of malaria parasites produced by the neighbor-joining method. Figures on the branches are bootstrap percentages based on 1000 replicates and only those 70 and above shown.

Vythilingam et al 2008  
Parasites & Vectors 1:26



Phylogenetic tree based on the non-repeat region of the circumsporozoite (csp) genes of malaria parasites produced by the Bayesian method. Figures on the branches are the posterior probabilities from the Bayesian analysis.

# DISCUSSION

- *Knowlesi* malaria occurs in this region (Jongwutiwes et al 2004, Zhu et al 2006, Cox –Singh et al 2008, Ng et al 2008, Luchavez et al 2008, Vythilingam et al 2008)
- With better molecular techniques one can differentiate between *P. malariae* and *P. knowlesi*

# DISCUSSION

- Chin *et al* 1968: Experimental transmission studies
- Due to declining anti-plasmodial immunity in humans
- Most cases occurring in malaria free areas not subject to control activities

# DISCUSSION

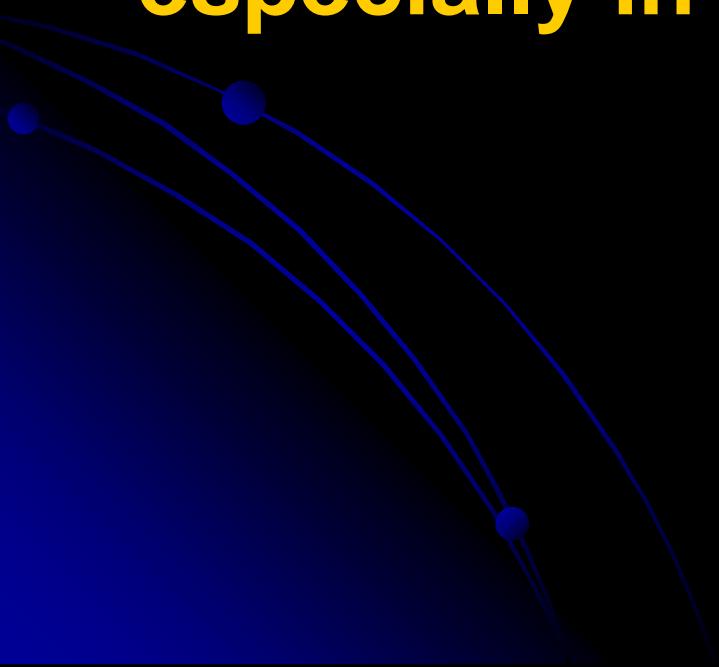
- This study has shown a link between *An. cracens*, humans and monkeys
- Early workers in Malaysia felt that simian malaria transmission to humans will be remote
- Development and deforestation- macaques close to human habitation

# DISCUSSION

- In the rain forests of Southeast Asia natural hosts of *P. knowlesi* – macaques abound so does the *An. leucosphyrus* group of mosquitoes
- Current control strategies are not going to work. Zoonotic transmission has to be widely publicized

# CONCLUSION

- Fifth human malaria parasite has been established and thus new strategies for malaria control should be considered especially in the elimination of malaria



# ACKNOWLEDGEMENTS

- Grant from the NIH, MOH Malaysia
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- Wildlife Department, Malaysia





**THANK YOU**