

ECOLOGY OF ARBOVIRUSES IN SRI LANKA: A SUMMARY OF STUDIES FROM 1984 -1990

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ABSTRACT

Studies on arboviruses carried out between 1984 -1990 at the University of Peradeniya are summarized here. The isolation of Nairobi sheep disease virus, Getah, Batai and Arkonam and serological evidence of the circulation of California encephalitis serogroup and Chandipura viruses have been reported for the first time in Sri Lanka. The ecology of Japanese encephalitis (JE) in areas with epidemic disease and sylvatic infection has been compared. A synchronous pattern of seroconversion in pigs and seroprevalence in cattle are more predictive of epidemic JE risk than overall porcine seroprevalence which is a good indicator of the presence of sylvatic JE rather than of human epidemic risk. Relatively modest changes in vector abundance associated with rainfall or agricultural practices have dramatic changes on the emergence of epidemic JE. Chandipura virus which is known to cause major outbreaks of encephalitis in India is endemic in Sri Lanka as are viruses belonging to the California encephalitis serogroup. Both these viruses are potential causes of human encephalitis and should be sought in patients with undiagnosed encephalitis. Nairobi sheep disease virus is endemic in animals (especially goats) and infects humans working closely with such livestock. Arboviruses remain an under-recognised cause of human and animal disease and an inter-sectoral multidisciplinary approach is needed to confront such emerging infectious disease threat.

Keywords: Japanese encephalitis, California serogroup, Alphavirus, mosquito, Chandipura virus

INTRODUCTION

Arboviruses (derived from arthropod-borne viruses) are a group of taxonomically diverse viruses that share a common mode of transmission between vertebrate hosts. The arthropods responsible for the transmission of these viruses are usually mosquitoes, ticks, sand flies, or less often, other insects. The majority of them fall into the taxonomic groups of the Togaviridae (Alphaviruses), Flaviviridae and Bunyaviridae. Most arboviruses (dengue and chikungunya being exceptions) are maintained in animal reservoirs with human infection being a tangential event, if at all, without much consequence to the survival of the virus. However, such infections can have significant implications for public health. Dengue continues to wreak havoc in many parts of Asia, Africa and South America (Messer *et al.*, 2002) while Japanese encephalitis (JE) remains a major cause of death and disability in Asia (Mackenzie *et al.*, 2002). West Nile virus has crossed the Atlantic to entrench itself in North America. Most intriguing of all, Chikungunya virus has re-emerged after a

“silence” of 40 years to become a major public health threat in many Asian countries and beyond (Powers and Logue, 2007).

A multi-disciplinary collaborative research program was initiated in 1983 by the late Prashantha Amerasinghe (Department of Zoology), Manel de S Wijesundera (Department of Parasitology) and J.S.M. Peiris (Department of Microbiology) at the University of Peradeniya to better understand the impact of ecological change on mosquito vectors of disease (Amerasinghe and Munasinghe, 1988; Amerasinghe *et al.*, 1991; Amerasinghe *et al.*, 1994) and to study the ecology of arboviruses in Sri Lanka. While some of the data on ecology of arboviruses has been published, other aspects of this work are so far only documented in M. Phil. and Ph.D. theses at the University of Peradeniya (Perera, 1986; Arunagiri, 1990). It is therefore opportune to summarize the findings of this research program on the ecology of arboviruses in Sri Lanka, particularly since some of these findings have

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increased in potential clinical relevance in the last decade. These studies have included a) Virus isolation from live caught mosquitoes and ticks, b) sero-epidemiological studies using neutralization tests and c) studies on the ecology and epidemiology of Japanese encephalitis virus in different regions of Sri Lanka.

Virus isolation from mosquitoes and ticks

Virus isolation was attempted from 178,181 un-engorged female mosquitoes collected between 1984-89 in the Anuradhapura (Dry zone), Dehiattakandiya (Intermediate zone area undergoing rapid ecological change as part of the Mahaweli System C irrigation program) and Kandy (Wet zone) areas (Peiris *et al.*, 1992; Peiris *et al.*, 1994). The viruses isolated are summarized in Table 1. Isolates of JE virus were obtained from different mosquito species and have helped define the mosquito species that transmit the virus in an epidemic setting. These studies also provided the first record of isolation of alphavirus Getah in the Indian subcontinent; the Bunyavirus Batai and the reovirus Arkonam from mosquitoes in Sri Lanka (Peiris *et al.*, 1994) and Nairobi sheep disease (NSD) virus from pools of *Haemaphysalis intermedia* ticks (Perera *et al.*, 1996) (Table 1).

Japanese encephalitis

Japanese encephalitis was endemic in Sri Lanka for decades (Hermon and Anandarajah, 1974; Vesenjak-Hirjan *et al.*, 1969) but the first major epidemic occurred in 1985 in the Anuradhapura district, North Central Province. There were substantial epidemics of varying

magnitude in the subsequent years. From 1987 – 1989, a prospective study of human disease, porcine seroconversion, vector abundance and virus carriage in mosquitoes was carried out in the Anuradhapura district. Based on JE specific IgM detection ELISA, 361 cases were virologically confirmed as JE between November 1987 and January 1988. Virus isolation incriminated the classical JE mosquito vectors *Culex tritaeniorhynchus*, *Culex gelidus* and *Culex fuscocephala* as vectors. These virus isolations were temporally correlated with porcine seroconversion and human disease (Peiris *et al.*, 1992). JE virus was also found in *Culex whitmorei* by virus isolation and in *Mansonia uniformis* by antigen detection ELISA (Peiris *et al.*, 1992; Peiris, 1994) but the overall vector abundance of these species was low and the role they played in the overall transmission of the virus is less clear. The vector responsible for the maintenance of the virus in its sylvan reservoir was not identified in these studies where mosquito trapping was done using light traps near pig pens. However, the isolation of JE virus from *Culex tritaeniorhynchus* and *Mansonia uniformis* in the forested / early settlement phase of the Mahaweli System C program at Dehiattakandiya and from *Culex pseudovishnui* in Kandy where there was no epidemic JE activity suggested that these species may be the responsible for the sylvatic transmission cycle. Both these species were found in substantial numbers (as many as 0.68 and 0.45 females / hr respectively) in the human biting collections in the forests of the Dry zone during the monsoon period (Amerasinghe and Munasinghe, 1988).

Table 1. Virus isolations from wild-caught female mosquitoes. (Tested in pools of up to 200 each)

Location	Viruses	Mosquito species (pools positive)	MIR*/1000 females
Anuradhapura 1986-89 n=59,422	Japanese encephalitis	<i>Culex tritaeniorhynchus</i> (4)	0.15
		<i>Culex gelidus</i> (5)	0.2
	Getah	<i>Culex tritaeniorhynchus</i> (6)	0.23
		<i>Culex gelidus</i> (1)	0.04
Arkonam	<i>Culex gelidus</i> (1)	0.04	
Dehiattakandiya 1985-89 n=62,079	Japanese encephalitis	<i>Culex tritaeniorhynchus</i> (3)	0.11
		<i>Mansonia uniformis</i> (2)	>1
	Arkonam	<i>Culex fuscocephala</i> (1)	0.13
Kandy 1984-87 n=56,680	Japanese encephalitis	<i>Culex pseudovishnui</i> (1)	0.16
	Getah	<i>Culex tritaeniorhynchus</i> (1)	0.09
		Batai	<i>Culex fuscocephala</i> (1)
		<i>Anopheles vagus</i> (1)	0.19

* MIR=Minimum infection rates / 1000 female mosquitoes tested.

Interestingly, we found evidence of JE seropositivity in persons engaged in slash and burn cultivation in the Dry zone forests (Perera, 1986). Ardeid water birds (e.g. *Bubulcus ibis*, *Ardeola greyii*) are the likely sylvatic reservoir hosts of JE and *Culex pseudovishnui* is likely to feed on such birds as well as on pigs and cattle and may potentially be the bridging vector between the sylvatic reservoir and the porcine amplifier host. The JE viruses isolated from *Cx.pseudovishnui* in Kandy and from a patient with JE encephalitis in Anuradhapura have been genetically sequenced and found to be closely related suggesting that the lack of epidemic JE in Kandy is not due to differences in virus virulence (Chen *et al.*, 1990) but is more likely due to abundance of vectors and amplifier hosts.

Porcine seroconversion followed a biphasic pattern, a slowly progressive phase of infection occurring through October 1987 when approximately 25% of the animals got infected, followed by a sharp synchronous burst of infection when the remaining animals got infected within a two week period in early November 1987. The human epidemic curve followed this burst of porcine seroconversion with a lag of 2-4 weeks and all ages of humans were equally affected, indicating an outbreak in an immunologically naive population. These findings confirmed the role played by pigs as an amplifier of the virus. Indeed it was highly likely that the reason why epidemic JE suddenly emerged in Anuradhapura, an ecologically stable rice growing region in 1985, was the introduction of small-holder pig-husbandry in the preceding few years to this ecosystem with high JE vector abundance because of intensive synchronized irrigated rice cultivation. The lessons learnt from this experience averted a similar mistake being repeated in the Mahaweli System C areas which were being developed for irrigated agriculture in the subsequent years.

Interestingly, there were few human cases in the subsequent year, *i.e.* 1988/9, viz. 11 confirmed cases in comparison to 361 cases in the preceding year. Thus, the difference of human disease burden in the two years was remarkable. This was not explained by build up of herd-immunity in the population since only 20% of the population had JE neutralizing antibody in early 1988. A vaccination campaign targeted at young children could not explain the marked reduction of disease in all ages. The notable difference in the two years was a roughly 5-fold reduction in the abundance of the major JE vectors, *Cx tritaeniorhynchus* and *Cx gelidus* (Peiris *et al.*,

1992). The change in vector abundance was attributable to changes in rainfall patterns in the two years and perhaps also to the disorganization of the irrigation water releases in the area due to the civil unrest that prevailed in the area. The correlation of rainfall pattern and JE disease deserves further study.

Interestingly, the change in vector abundance was associated with a subtle change in the profile of porcine seroconversion. While the overall porcine seroprevalence reached almost 100% in both years, in 1998 (low human JE year) there was an absence of the sharp synchronous burst of porcine seroconversion observed in 1987. This suggested that it was not the overall porcine seroprevalence that predicted high JE risk but rather the tightly synchronized seroconversion in pigs. These hypotheses were confirmed by ecological studies of JE in different agroclimatological regions of Sri Lanka. Porcine seroprevalence did not correlate with risk of human JE; for example Kandy had a cumulative or cross sectional porcine seroprevalence not dissimilar to that of Anuradhapura. What did differ was the lack of synchronized seroconversion over a short period of time. Indeed, our studies suggested that bovine seroprevalence correlated better with human epidemic JE risk than does porcine seroprevalence (summarized in Table 2), probably because bovines were a spillover host for JE virus just as humans are. On the other hand, cross sectional porcine seroprevalence is a very sensitive indication of JE virus activity in the sylvatic reservoir but not necessarily a predictor of human spillover infection (Peiris *et al.*, 1993a).

Dengue viruses

Uncomplicated dengue fever was endemic in Sri Lanka for decades (Maheswaran *et al.*, 1976) but the emergence of dengue haemorrhagic fever as a major public health threat in Sri Lanka was a more recent phenomenon (since 1989) and occurred subsequent to the period covered by the studies reported here. As dengue was not a particular focus of our studies the mosquito collection methods were not targeted the classical dengue vectors *Aedes aegypti* and *Aedes albopictus*. Seroprevalence (neutralizing antibody) to dengue 2 virus in humans in Kandy, Anuradhapura and in prior settlers involved in slash and burn agriculture in the Maduru Oya forested environment was 16%, 25% and 21% respectively, indicating that dengue was widespread in the island even pre-1988 although dengue haemorrhagic fever was then uncommon (Perera, 1986).

Interestingly, a seroepidemiological study of torque macaques (*Macaca sinica*) in Polonnaruwa in 1987 revealed evidence of high (94%) seroprevalence to dengue 2 (but not to JE virus) (Peiris *et al.*, 1993b). A follow-up study conducted in 1995 by De Silva *et al.*, (1999) confirmed that the dengue epizootic was limited in both space and time, indicting that this was most likely a spill over of infection from humans to macaques rather than a result of endemic virus transmission within this ecosystem

Alphaviruses: Getah and chikungunya

In agreement with studies in Malaysia, we found *Cx tritaeniorhynchus* to be the likely vector of Getah in Sri Lanka. Sero-epidemiological studies using micro-neutralization tests using one of these isolates (An-432) showed that pigs had the highest sero-prevalence while cattle, goats, sheep, dogs, chickens and ducks also have evidence of infection. Human sero-prevalence was low (0.6%) in most parts of the country (Table 3) (Arunagiri, 1990). Chikungunya virus is a related alphavirus that caused major disease outbreaks in Sri Lanka in the 1960's (Hermon,

1967). There was no detectable antibody in sera from 321 children aged 5-15 and very low seroprevalence (1/180) in adults. This single seropositive serum specimen had low neutralizing antibody titre to chikungunya (1/40) and Sindbis (1/20). Overall the studies confirmed that chikungunya was not endemic in Sri Lanka in the last decades and indicated that the population was largely susceptible to the Chikungunya virus and also to other alphaviruses (Table 4) (Arunagiri, 1990). This is relevant as baseline information in view of the major outbreaks of chikungunya fever that have erupted in Sri Lanka, India and other parts of Asia in recent years (Powers *et al.*, 2007).

Batai and Arkonam virus

The bunyamwera serogroup virus Batai was found to be endemic in Sri Lanka by both virus isolation and by sero-epidemiological studies (Peiris *et al.*, 1994; Arunagiri, 1990). Antibody to Batai virus was common (20-30%) in cattle, sheep and goats but human seroprevalence was low (Arunagiri, 1990).

Table 2. Correlation of Human Japanese Encephalitis Case Notifications with Serological and Entomological Parameters. (Source: Peiris, 1996)

	<u>Anuradhapura</u>	<u>Ragama</u>	<u>Kandy</u>	<u>High elevation / Wet zone</u>
Elevation (m)	150	30	530	1270-2070
JE case notifications / 100,000 (1990)	26	4-72	0.1	0
Human JE sero-prevalence (school age)	17%	28%	3%	2%
Porcine JE sero-prevalence (>13 months)	89%	71%	76%	17%
Pattern of porcine sero-conversion	Synchronous	Synchronous	Asynchronous	Asynchronous
Cattle JE sero-prevalence	45%	71%	4%	0%
<u>Mosquito abundance(per trap night)</u>				
<i>Cx. tritaeniorhynchus</i>	898	28	49	1.3
<i>Cx. gelidus</i>	433	96	37	3.2
<i>Cx. fuscocephala</i>	77	10	65	9.8

Table 3. Seroprevalence of neutralizing antibody to Getah virus (An432). (Source: Arunagiri, 1990)

Agro-climatic zone	Human (5-15 yr)	Pigs	Cattle	Goat	Dog	Chicken
Dry Zone (NCP)	0/152 (0%)	75/181 (41%)	13/75 (17%)	12/151 (8%)	2/16 (13%)	8/93 (9%)
Dry Zone (MSC)	1/142 (1%)	5/13 (38%)	2/18 (11%)	ND	ND	ND
Wet Zone <1000 ft	ND	17/228 (7%)	8/35 (23%)	2/37 (5%)	ND	ND
Wet Zone 1000-3000 ft	2/229 (1%)	2/161 (1%)	22/136 (16%)	4/62 (6%)	0/12 (0%)	0/28 (0%)
Wet Zone >3000 ft	0/179 (0%)	24/338 (7%)	0/23 (0%)	ND	ND	ND

Abbreviations: NCP=North Central Province; MSC=Mahaweli System C; ND=Not done

Arkonam is an Orbivirus within the family Reoviridae. It has been previously detected in India but its importance in relation to human or animal disease was unknown. While no seroprevalence studies were done in Sri Lanka so far, studies in India suggest that human infection is not uncommon (Dandawate and Shope, 1975).

Nairobi sheep disease virus

Nairobi sheep disease (NSD) virus was isolated from pools of *Haemaphysalis intermedia* ticks collected from the goat breeding station at Kotukachchiya in the North Western Province of Sri Lanka (Perera *et al.*, 1996). This was the first record of its isolation in Sri Lanka although Ganjam virus which is now recognized to be indistinguishable from NSD virus has been isolated from sheep in association with febrile and sometimes fatal illness in India with symptoms similar to those of NSD in East Africa (Ghalsasi *et al.*, 1981). Seroepidemiological studies indicated that human and goat exposure is not uncommon on the farm we studied. Its relation to human and goat disease deserves further investigation.

California serogroup viruses

Sero-epidemiological studies revealed the activity of a number of other arboviruses in Sri

Lanka. The most important of these was the conclusive documentation of the presence of California serogroup Bunyaviruses and Chandipura virus for the first time in Sri Lanka. Bardos and colleagues had previously reported inconclusive evidence of antibody to Tahyna virus in the animal sera from Sri Lanka (Bardos *et al.*, 1983). Both California serogroup and Chandipura viruses are known causes of viral encephalitis in humans. Neutralization tests revealed the presence of a virus closely related to Snowshoe hare virus and another related to Melao virus (Arunagiri *et al.*, 1991). In a comparative study of sera from school age children (8-16 years) in different agro-climatological regions of Sri Lanka, the highest sero-prevalence (6%) in children being found in those living within the dry zone forested areas of the country with 1% seroprevalence in children elsewhere. Pigs, cattle, dogs, rodents and chicken had significant sero-prevalence to these viruses (Arunagiri *et al.*, 1991). Toque macaques (*Macaca sinica*) in a sylvatic habitat in the North Central Province of Sri Lanka had high seroprevalence to California serogroup viruses, with an age depended increase in seroprevalence suggesting endemic virus transmission (Peiris *et al.*, 1993).

Table 4. Seroprevalence of neutralizing antibody to Chikungunya and Sindbis viruses. (Source: Arunagiri, 1990)

	Chikungunya	Sindbis
Adult sera	1/180 (0.6%)	1/298 (0.34%)
Children (5-15 years)	0/321 (0%)	1/485 (0.2%)

Sera collected from different agro-climatic regions of Sri Lanka (see Table 2). There was no obvious difference in seroprevalence when stratified by geographical region.

Table 5. Seroprevalence of neutralizing antibody to Chandipura virus. (Source: Arunagiri, 1990)

Agro-climatic zone	Human (5-15 yr)	Pigs	Cattle	Goat
Dry Zone (NCP)	1/103 (1%)	17/137 (12%)	27/75 (36%)	8/105 (7.6%)
Dry Zone (MSC)	8/142 (5.6%)	3/7 (42%)	17/18 (94%)	ND
Wet Zone <1000 ft	ND	3/101 (3%)	18/32 (56%)	2/34 (5.9%)
Wet Zone 1000-3000 ft	2/229 (0.9%)	4/114 (3.5%)	57/136 (42%)	4/62 (6.5%)
Wet Zone >3000 ft	3/177 (1.7%)	4/159 (2.5%)	5/23 (22%)	ND

Abbreviations: NCP=North Central Province; MSC=Mahaweli System C; ND=Not done

Table 6. Arboviruses documented in Sri Lanka.

Virus	Isolated from humans in Sri Lanka	Human sero-positivity in Sri Lanka	Animal sero-positivity in Sri Lanka	Virus isolated from vector in Sri Lanka	Human disease world-wide	
					Human disease	CNS disease
Japanese encephalitis	+	+	+	+ ¹	+(SL) ³	+(SL)
Dengue types 1-4	+	+	+ ¹ (<i>Macaques</i>)	-	+(SL)	+
Chikungunya	+	+	?	-	+(SL)	-
California encephalitis serogroup	-	+ ¹	+ ¹	-	+	+
Chandipura	-	+ ¹	+ ¹	-	+	+
Nairobi sheep disease	-	+ ¹	+ ¹	+ ¹	+	+/-
Bhanja	-	+ ¹	+	-	+	+
Getah	-	+ ¹	+ ¹	+ ¹	-	-
Batai	-	+ ¹	+	+ ¹	?	-
Arkonam	-	-	-	+ ¹	-	-
Sathuperi	-	+ ¹	+ ¹	-	-	-
Wanowrie	+ ²	-	-	-	+	+

¹ First documented during following studies (Peiris *et al.*, 1994; Arunagiri *et al.*, 1991; Arunagiri, 1990, Perera, 1996).

² Pavri *et al.*, 1976 ³ SL = human disease documented in Sri Lanka.

Table is reproduced with modification from Peiris, 1996.

California serogroup viruses have been reported to circulate in Azerbaijan and Tajikistan in Central Asia (Lvov *et al.*, 1980) and human infection documented in Shanghai (Gu *et al.*, 1984). Since these viruses are well established causes of human encephalitis in North America and febrile disease in Europe (Mertz, 2002), it is important that their role in undiagnosed viral encephalitis in Sri Lanka is investigated.

Chandipura virus

Chandipura virus is a Vesiculovirus in the family Rhabdoviridae, typically transmitted by Phlebotomus sand flies. Seroprevalence of Chandipura virus was consistently found to be highest in the Mahaweli System C areas, whether in cattle, pigs or humans (Table 5). Three of 80 individuals recently settled in the Mahaweli System C area seroconverted to the virus in their first year of residence (Arunagiri, 1990). Cattle had the highest sero-prevalence of antibodies to Chandipura virus with seroprevalence as high as 94% in the Dry Zone Mahaweli System C areas but virus activity being present even at higher altitudes of >3000 ft. Pig seroprevalence was 42% in the Mahaweli System C area and lower at high altitudes. At the time these studies were carried out, the clinical

importance of Chandipura virus was less well appreciated but it has more recently been incriminated as a cause of epidemic encephalitis in Andhra Pradesh, India, with some outbreaks affecting over 300 children (Rao *et al.*, 2004). The disease starts as an influenza like illness, associated with abdominal pain, vomiting and altered consciousness and impaired neurologic function. Chandipura virus should be particularly investigated for in patients with viral encephalitis or encephalopathy in the Dry zone areas of Sri Lanka.

CONCLUSION

The arboviruses documented in Sri Lanka in our studies as well as by others are summarized in Table 6. Arbovirus infections remain the archetypal ecological disease where an integrated effort of those involved in public health, entomology, animal health, agriculture and wildlife needs to be harnessed. The late Prashantha Amerasinghe was an excellent example of one who could bridge these inter-sectoral divides and we remain indebted to him for playing a key role in these studies, among

many other research studies he carried out during an exceptionally productive scientific career. The need for such an inter-sectoral approach is highlighted also in more recent emerging infectious disease problems such as avian influenza, Nipah and SARS.

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