

Are Haematophagous Insects Vectors for HTLV-I?

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A. Introduction

Human T-lymphotropic virus I (HTLV-I) and its associated malignancy, adult T-cell leukaemia [19, 30, 31, 37] are endemic in southern Japan [18, 26], the Caribbean Basin and neighbouring mainland America [1, 6, 10, 14, 23], and sub-Saharan Africa [5, 20, 38].

The natural history and mode of transmission of HTLV-I infection is largely unknown. Worldwide, the infection as indicated by the presence of antibodies is confined mostly to tropical areas of high humidity in Japan [36], the Caribbean and tropical South America [7, 23], and Central Africa [4, 5, 20, 38]. Sero-positivity for HTLV-I and the associated risk of adult T-cell leukaemia/lymphoma has been reported among immigrants to temperate regions from the Caribbean [10, 15]. More recently, southern Italy [22] and the Arctic regions [32] have been identified as possibly additional endemic regions. In the Caribbean, sero-positivity has been found almost entirely in people of African or part-African descent, although there are recent reports of HTLV-I-positive adults among Amerindians in Cayenne [11] and Venezuela [23] and Campuchian immigrants to Cayenne [11].

Earlier studies, especially in Japan, have suggested that HTLV-I might be transmitted by sexual contact [36], from mother to

child via transplacental passage [21] or breast feeding [28], by blood transfusion [16], or by insect vectors [35]. Sexual transmission has been emphasised as a possible major route, but it cannot easily explain the demographic distribution of sero-positivity or the relatively high incidence of HTLV-I antibodies in children whose parents are sero-positive [17]. Neither does it adequately explain why the highest rates of sero-positivity should be observed in 70-year-old Japanese women.

B. A Role for Haematophagous Insects?

The possibility has been raised that Africans represent the natural host of HTLV-I, and that the virus was introduced in this way to the West Indies and perhaps also to Japan [13, 20]. In Britain, HTLV-I sero-positivity is found in first generation but very rarely in second generation West Indian people of African descent (Table 1) [15] (M. F. Greaves, T. A. Lister, S. Pegram and L. Chan, unpublished observations). Thus, the circumstances required for transmission may be largely confined to tropical and subtropical areas of high humidity, and – strikingly – to some Arctic regions [32]. Within endemic areas strong associations exist between sero-positivity to HTLV-I and evidence of exposure to arthropod-transmitted diseases such as filariasis [35], malaria [3] and equine encephalitis [23]. These relations might imply human transmission of HTLV-I by mosquitoes, shared environmental factors which promote the transmission of HTLV-I and arthropod-borne disease, or

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Table 1. HTLV-I antibodies in United Kingdom residents of Caribbean origin

1. Relatives of HTLV-I ⁺ adult T-cell leukaemia patients:
23 Relatives of 6 patients
Sero-positive: 0/12 UK born (of Afro-Caribbean parentage)
4/11 Caribbean born
2. Non-leukaemic serum donors (unrelated to ATL patients):
Series 1: Hospital out-patients
Sero-positive: 6/70 – all positives were born in Caribbean [15]
Series 2: Hospital out-patients + normals
Sero-positive: 0/70 UK born (of Afro-Caribbean parentage)
6/130 Caribbean born

potentiation of the response to one infection by previous exposure to the other [3].

Involvement of mosquitoes or other haematophagous insects in the transmission of HTLV-I would go some way towards explaining the endemicity of this virus in Arctic regions where such insects are exceedingly common. Recent collaborative studies we have conducted on the Caribbean island of Trinidad provide, we believe, further support for the idea that insects might be vectors for HTLV-1 [25]. Between 1977 and 1981, blood samples were obtained for a cardiovascular survey of all adults aged 35–69 years and resident within a geographically defined area of Port-of-Spain, Trinidad [24]. In most respondents, sufficient serum had been stored for screening of the community for evidence of HTLV-I infection. As far as we are aware, this is the first systematic study of the distribution of HTLV-I seropositivity in a total community.

The population of Trinidad is of African, Indian, European, Chinese, Lebanese and Syrian descent. The forbears of those of African origin arrived after 1776, coming either directly from West Africa or by way of neighbouring islands such as Grenada. People of Indian descent came from 1845 onwards, mostly from northern India. Smaller numbers of Chinese entered Trinidad after 1852.

Port-of-Spain lies on the coast of N.W. Trinidad and has a humid tropical climate. The survey was conducted in a defined area encompassing contiguous sectors of two suburbs of the city which we refer to as sectors A and B. Although in general the A-sector community is of lower socio-economic standing than sector B, there is no appreciable difference between Africans and Indians in living standards within the A sector.

Sera were screened for HTLV-I with an enzyme-linked immunosorbent assay (ELISA) [33]. Samples positive with this test were examined with an ELISA modification of a previously described competitive assay [40]. Only subjects whose sera were positive to both assays were considered sero-positive for HTLV-I.

Details of this study are published elsewhere [25]. Several important facts emerged: 1. Individuals of Asian (Indian) descent were infected at a lower rate (1.4%) than Afro-Caribbean blacks (7.0%). 2. As in Japan, sero-positivity rates were age and sex associated, the highest rates (12.3%) being observed in females over 65 years of age. 3. Black males living in the less-prosperous sector A were infected at a higher rate (5.4%) than those living in sector B (2.6%). 4. Sero-positivity was significantly ($P < 0.001$) associated with poor-quality housing. 5. Sero-positivity was significantly ($P < 0.025$) associated with living distances of less than 30 m from open water courses.

We have suggested [25] that, when taken together, the household clustering of HTLV-I infection (documented in Japan), the increased risk to adult females, the association with poor-quality housing and proximity to insect breeding sites (i.e. open, stagnant water) implicate an insect vector of intensively domestic habit. One candidate would be the mosquito species *Aedes aegypti*, which is known to have limited dispersal from its breeding grounds [27], but other domestic insects including mites and ticks would also be candidates. Women might be more likely to be infected than their husbands by virtue of their spending more time in the house and therefore incurring a greater risk of exposure. In the Caribbean region, non-black ethnic groups might be infected only by co-habitation with people of African descent who probably provide the

host reservoir for the virus, and this would need to be prolonged if infectivity were very low. This might then explain why Indians, though infected with HTLV-I, have a much lower sero-positivity rate in Trinidad than do inhabitants of African origin.

The increasing sero-positivity with age still requires an explanation. One possibility is that it reflects the requirement for continuous exposure to potential sources of virus coupled with inefficient transmission. Another possibility is that the age association is merely a mirror of historical events, with young people now being infected at a much lower rate due, for example, to improving living conditions. We have assumed that if blood-sucking insects are capable of transmitting HTLV-I this will be via the purely mechanical transfer of infected lymphocytes. At this stage, however, it remains possible, though perhaps less likely, that insects serve as a biological vector supporting viral replication, as happens, for example, with the Japanese encephalitis arbovirus [34]. This is currently being investigated.

Finally, it is important in this context to re-emphasise the similarity between HTLV-I and bovine leukaemia virus (BLV) with respect to viral structure and disease similarities [8, 39]. Although BLV in developed countries may commonly be transmitted iatrogenically, there may well be a role for haematophagous insects as mechanical vectors, especially in tropical regions. Experimental evidence for this has been produced [2, 9, 12]. BLV does not, however, spread readily within a herd of cows, and physical contact is almost certainly required [29]. This might again implicate an insect vector other than common mosquitoes.

It is of interest that BLV can be experimentally transferred by injecting a sero-negative cow with 100 μ l of blood or around 1000–2000 lymphocytes from an infected cow, but that this value drops by 3 orders of magnitude if the donor cow has a BLV-associated lymphocytosis (A. Burny, this volume). The latter corresponds to the transfer of less than 0.1% of the volume of a mosquito blood feed which could easily be carried on the mouthparts. Perhaps HTLV-I is effectively transmitted only by blood-sucking insects that have had an interrupted meal on a sero-positive individual with some de-

gree of lymphocytosis. This might then explain the apparent need for close familial contact over a prolonged period.

It is clear that HTLV-I has been transmitted by blood transfusion in Japan [16], and we do not contest that infection can be spread via breast feeding or sexual contact. Although several modes of transmission may be possible, we hypothesise that insect vectors play a more significant role than has hitherto been appreciated.

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