

**Table 2.11. Case-control studies of HTLV and co-factors *strongyloides stercoralis* (Ss)**

Reference, study location and period	Characteristics of cases	Characteristics of controls	Detection method	Exposure categories	No. of exposed cases	Relative risk (95% CI)	Adjusted potential confounders	Comments
Plumelle et al. (1997) Martinique 1983–1995 cross-sectional analysis of case series	19 ATLL cases, Ss+ ages 23–57 yrs at dx, 11 men & 8 women	19 ATLL cases, Ss- ages 31–90 yrs at dx, 9 men & 10 women	Anti-HTLV-1 by ELISA with Western blot; Ss by Baerman method					<u>Age at Dx:</u> Ss+ pts were younger at dx ( $p=0.0002$ )
Gabet et al. (2000) French Guyana, French West Indies. Cross-sectional analysis	18 HTLV-1 carriers co-infected with Ss: 8 from West Guyana, 10 from Martinique; 11 males; aged 10–72 yrs	15 HTLV-1 carriers not infected with Ss: 8 from West Guyana; 7 from Martinique; 6 males; aged 30–54 yrs	Proviral load by quantitative PCR; clonal expansion by semi-quantitative PCR of flanking sequences. Ss by Baerman method	HTLV-1 clonal frequency: $\geq 1/300$ ; $\geq 1/1500$ – $< 1/300$ ; $\geq 1/3000$ – $< 1/1500$ PBMCs			<u>Proviral load</u> Ss+ carriers had higher proviral loads ( $p=0.009$ ) <u>Clonality</u> Ss+ carriers had more HTLV-I+ clones ( $p=0.024$ )	
Satoh <i>et al.</i> (2002) Japan, Okinawa	31 HTLV-1 carriers co-infected with Ss	10 HTLV-1 carriers not infected with Ss	Anti-HTLV-1: PAA and indirect immunofluorescence; proviral load by real-time quantitative PCR; clonal expansion by inverse-long PCR; Ss by fecal culture and PA assay				<u>Median proviral load</u> Ss+ = 15.3% v. 3.9% ( $p<0.005$ )	