Detection of Anti–HTLV-I Tax Antibodies in HTLV-I Enzyme-Linked Immunosorbent Assay–Negative Individuals


The HTLV-I tax gene protein (Tax) is not packaged within the mature viral particle from which the proteins for the commercially available enzyme-linked immunosorbent assay (ELISA) are derived. Screening of 182 individuals within a cohort of white intravenous (IV) drug abusers, previously identified as having an increased incidence of HTLV-I infection, demonstrated that seven of them had antibodies to the HTLV-I Tax protein but tested negative in HTLV-I ELISAs and Western blots prepared from purified virion proteins. Three out of 35 individuals in other behaviorally defined high-risk groups also displayed this limited pattern of reactivity to HTLV-I proteins. The presence of the anti-HTLV-I p40/Tax antibodies was determined by radioimmunoprecipitation assay (RIPA), which also revealed low levels of anti-env reactivity. The specificity of the anti-p40 reactivity was confirmed on specific Tax ELISAs and Western blots prepared from recombinantly produced Tax. In vitro gene amplification by the polymerase chain reaction (PCR) was used to establish the presence of sequences homologous to HTLV-I proviral DNA in four/four of these HTLV-I ELISA negative, Tax ELISA/Tax western blot/RIPA positive individuals. These data suggest that the true incidence of HTLV-I infection within high-risk cohorts is greater than previously reported.

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formed as previously described, except that in some cases two identical mixed cultures of SLB-I and MT2 cells were metabolically labeled with \(^{35}\)S methionine and \(^{35}\)S cysteine, respectively, for four hours. These two cultures were then combined to prepare a doubly labeled mixed lysate to ensure high levels of all HTLV-I proteins. HIV-1 ELISA and HTLV-I ELISA and Western blot kits were provided by Cellular Products Inc (Buffalo, NY) and used according to the manufacturer's specifications. HTLV-II ELISAs were performed exactly as described, except that HTLV-II virion proteins from banded particles produced by the MOT cell line were used as the source of antigen. The HTLV-I tax protein ELISA has been previously described. Briefly, assays were performed by incubating 1:20 dilutions of serum samples for one hour at 37°C in duplicate microtiter wells coated with 200 ng of purified recombinant HTLV-I tax protein. After rinsing with phosphate-buffered saline (PBS)/0.05% Tween 20, bound human antibody was detected with protein A conjugated to horseradish peroxidase and o-phenylenediamine/H_2O_2. The reaction was measured at 490 nm with a Dynatech MR700 microplate reader. Tax Western blots were prepared by electrophoresing purified recombinant tax protein through a Laemmli gel system and transferring the protein to nitrocellulose by electroblotting. The filter was washed for 15 minutes in 1X PBS (GIBCO, Grand Island, NY), 0.3% Tween 20, and stored on 3MM Whatman paper in the dark. Strips containing 70 ng of protein were cut from the filter (50-mm length x 3-mm width). Individual strips were placed in polypropylene tubes and blocked for one hour with 1X PBS, 0.3% Tween 20, 3% bovine serum albumin (BSA) on a rocking platform at 24°C, followed by washing three times with 1X PBS, 0.3% Tween 20. Serum samples were diluted 1:50 in wash solution, added to the filter strips, and incubated overnight on a shaker at 4°C. Filters were then washed three times as above. A 1/1,000 dilution of an antihuman IgG conjugated to horseradish peroxidase was incubated with each strip at 24°C on a rocker for one hour. Filters were washed three times, substrate (4-chloro-1-naphthol) was added, and incubation for color development was carried out for 20 minutes in the dark at 24°C on a shaker. Substrate was removed, and the filter was washed with distilled water.

**Gene amplification analyses.** PCR was performed in a Perkin-Elmer Cetus Thermal Cycler (Norwalk, CT) with the thermostable DNA polymerase, Taq, isolated from *Thermus aquaticus* for 45 cycles as described. Amplified DNAs were assayed in spot-blot and liquid hybridization formats as described. The regions of the HTLV-I tax and pol chosen for amplification and detection have been previously described.

### RESULTS

The sensitivity and specificity of the HTLV-I ELISA, and Western blot assays prepared from viral proteins, and the HTLV-I RIPA based on infected cell lysates have been previously established (Poiesz et al; Anderson et al; Slamon et al, manuscripts in preparation). The validity of the recombinant Tax ELISA and a cut-off determination were established by running 187 normal human serum samples and measuring the resultant absorbances. The mean of these samples was determined to be 0.126 with an SD of 0.073. Using the same convention that we employ for the HTLV-I virion protein ELISA, where the cut off is defined as the mean of the normals plus 5 SDs, a cut-off value of \(0.500\) was established. The specificity and sensitivity of the Tax ELISA were further investigated in a double-blind analysis in which samples from SUNY HSC were coded and sent to Triton Biosciences Inc. The mean absorbance readings for various groups were as follows: ATL and HTLV-I associated myelopathy (HAM) patients \(2,000\) (four had absorbance readings between 1.523 and 1.997; 16 were >2.000), \(n = 20\); random Red Cross blood donors 0.155, \(n = 60\); other diseases, including acquired immunodeficiency syndrome (AIDS), lymphoma, granulomatous syndromes, autoimmune diseases 0.187, \(n = 90\). A single patient with an HTLV-I benign lymphoproliferative disease gave an absorbance reading of 1.090. These findings established the sensitivity and specificity of the Tax ELISA to be 100% and 99.1% (3/337 normals and individuals with other diseases tested above the cutoff), respectively. All patients with clinically expressed HTLV-I disease tested in this study were positive in the Tax ELISA.

The HTLV-I virion protein ELISA was the least sensitive assay of these methods for the detection of immunoreactivity directed against the HTLV-I tax gene protein (Table 1). The HTLV-I RIPA was approximately 50 times more sensitive, and the specific assays (ELISA and Western blot using recombinantly prepared Tax) designed to specifically detect seroreactivity to Tax were approximately three orders of magnitude better than the standard HTLV-I ELISA. A

### Table 1. Comparison of Various Serologic Assays for Their Ability to Detect Antibodies Against HTLV-I Viral Proteins and Tax

<table>
<thead>
<tr>
<th>Assay Serum</th>
<th>HTLV-I RIPA* 40-Kd Tax</th>
<th>ELISA†</th>
<th>Tax‡ Western</th>
<th>HTLV-I† ELISA</th>
<th>HTLV-II‡ ELISA</th>
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</thead>
<tbody>
<tr>
<td>αTax rabbit</td>
<td>5,000</td>
<td>90,000</td>
<td>&gt;100,000</td>
<td>90</td>
<td>500</td>
</tr>
<tr>
<td>Normal rabbit</td>
<td>0</td>
<td>500</td>
<td>0</td>
<td>0</td>
<td>ND</td>
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<tr>
<td>αHTLV-I rabbit</td>
<td>0</td>
<td>500</td>
<td>9,000</td>
<td>ND</td>
<td></td>
</tr>
<tr>
<td>HAM patient</td>
<td>1,000</td>
<td>700</td>
<td>8,000</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Laboratory worker</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td>0</td>
<td></td>
</tr>
</tbody>
</table>

All data are given as the reciprocals of the highest dilution at which the serum gave a positive signal.

Abbreviation: ND, not determined.

*End points for RIPAs were determined by assessing the highest serum dilution at which a 40-Kd band could be visualized by fluorography. Sera were run at \(2 \times 10^{-2}, 10^{-2}, 2 \times 10^{-2}, 10^{-3}, 10^{-4},\) and \(10^{-5}\) dilutions.

†End points for ELISAs were determined by plotting the OD_max results vs serum dilution and determining where the individual curves crossed the cut-off line, defined as 5 SDs above the mean of the normal control samples. Sera were run at \(2 \times 10^{-2}, 10^{-2}, 2 \times 10^{-2}, 10^{-3}, 10^{-4},\) and \(10^{-5}\) for the tax protein ELISA and at \(2 \times 10^{-3}, 10^{-3}, 2 \times 10^{-3}, 10^{-4},\) and \(10^{-5}\) for the HTLV-I ELISA.

‡End points for all Western blots were determined by assessing the highest serum dilution at which a 40-Kd band could be visualized following color development. Sera were run at \(2 \times 10^{-3}, 10^{-3}, 2 \times 10^{-3}, 10^{-4},\) and \(10^{-5}\) dilutions.
Tables 2. Serum Reactivities of Retroviral Risk-Group Members  

<table>
<thead>
<tr>
<th>Patient</th>
<th>HTLV-I p40 Band</th>
<th>HTLV-I ELISA</th>
<th>HTLV-I Western</th>
<th>Tax ELISA</th>
<th>Tax Western</th>
<th>HTLV-II ELISA</th>
<th>HIV-I ELISA</th>
<th>Risk Group</th>
<th>Race</th>
</tr>
</thead>
<tbody>
<tr>
<td>5044</td>
<td>+</td>
<td>19 (-)</td>
<td>-</td>
<td>952 (+)</td>
<td>+</td>
<td>-</td>
<td>-</td>
<td>IVDA</td>
<td>W</td>
</tr>
<tr>
<td>5048</td>
<td>+</td>
<td>177 (-)</td>
<td>-</td>
<td>&gt;2,000 (+)</td>
<td>+</td>
<td>-</td>
<td>+</td>
<td>IVDA</td>
<td>W</td>
</tr>
<tr>
<td>5079</td>
<td>+</td>
<td>88 (-)</td>
<td>-</td>
<td>727 (+)</td>
<td>+</td>
<td>-</td>
<td>+</td>
<td>IVDA</td>
<td>W</td>
</tr>
<tr>
<td>5173</td>
<td>+</td>
<td>36 (-)</td>
<td>-</td>
<td>1,308 (+)</td>
<td>+</td>
<td>-</td>
<td>-</td>
<td>IVDA</td>
<td>W</td>
</tr>
<tr>
<td>5324</td>
<td>+</td>
<td>142 (-)</td>
<td>-</td>
<td>998 (+)</td>
<td>+</td>
<td>-</td>
<td>+</td>
<td>IVDA</td>
<td>B</td>
</tr>
<tr>
<td>5327</td>
<td>+</td>
<td>32 (-)</td>
<td>-</td>
<td>1,714 (+)</td>
<td>+/-</td>
<td>-</td>
<td>-</td>
<td>IVDA</td>
<td>W</td>
</tr>
<tr>
<td>5334</td>
<td>+</td>
<td>49 (-)</td>
<td>-</td>
<td>1,564 (+)</td>
<td>+</td>
<td>-</td>
<td>ND</td>
<td>MA</td>
<td>W</td>
</tr>
<tr>
<td>5351</td>
<td>+</td>
<td>56 (-)</td>
<td>ND</td>
<td>533 (+)</td>
<td>+</td>
<td>-</td>
<td>+</td>
<td>IVDA</td>
<td>W</td>
</tr>
<tr>
<td>5354</td>
<td>+</td>
<td>53 (-)</td>
<td>ND</td>
<td>393 (+/-)</td>
<td>+</td>
<td>+</td>
<td>+</td>
<td>HS</td>
<td>W</td>
</tr>
<tr>
<td>5370</td>
<td>+</td>
<td>32 (-)</td>
<td>ND</td>
<td>314 (+/-)</td>
<td>+</td>
<td>-</td>
<td>-</td>
<td>F1</td>
<td>B</td>
</tr>
</tbody>
</table>

All serum samples analyzed were identified in a HTLV-I RIPA as having antibodies against a 40-Kd protein present in cellular lysates prepared from SLB-I.

Abbreviations: MA, malarial Africa; HS, homosexual; F1, offspring of risk group member; W, white; B, black; ND, not determined.
positive. Analyses of these patients’ DNAs after amplification for HTLV-1 pol sequences showed that DNA from three of these four patients contained HTLV-I sequences as determined by both liquid and spot-blot hybridization analyses (Fig 3C). The single pol-negative DNA (5327) gave the strongest signal with the tax primer pair.

DISCUSSION

A serologic analysis had been previously conducted to determine the extent of exposure to HTLV-I among a cohort of primarily white, US born IVDA's from the New York City area. Six of 162 (3.7%) patients, all white, were found to be repeatedly reactive in an HTLV-I ELISA prepared from viral proteins. All of these HTLV-I–positive results were confirmed by RIPA and/or Western blot as having antibodies directed against HTLV-I gag, env, and tax gene products. In the present study we describe a phenomenon of limited reactivity to HTLV-I proteins in 7/162 individuals from this IVDA cohort and 3/35 individuals from other risk groups. When these results are combined with the initial survey, it brings the total number of HTLV-I seroreactive individuals in the IVDA cohort to 13/162 (8.0%), thereby more than doubling the number of suspected HTLV-I–infected individuals. These ten patients whose sera were negative in a commercial HTLV-I ELISA displayed strong anti-Tax and moderate anti-env reactivity by RIPA but very little if any anti-gag reactivity. These findings, when combined with the knowledge that shearing of the HTLV-I envelope proteins occurs during gradient purification, indicate that the ELISAs presently available contain predominantly gag-encoded epitopes with lower levels of env proteins and almost no Tax.

The detection of HTLV-I homologous sequences in the DNA of all four of the RIPA+ /ELISA– individuals from whom cells were available suggests that a majority of the patients identified by RIPA as having predominantly anti-tax antibodies also carry the HTLV-I provirus. The provirus from the patient who was LH negative for tax probably contains a small mutation that affects oligonucleotide hybridization, but the seroreactivity data indicate that the gene is transcribed. The patient (5044) displaying the variable tax gene LH results between samples represents an interesting case. Reamplification and detection of both samples with both assay formats confirmed the initial observa-
tions. These findings suggest the possibility of infection with multiple strains, or strain evolution, both phenomena which have been documented with HIV-1. Future work will involve cloning and sequence determination of each of these samples. The region of pol amplified by SK54/55 may represent a variable region within HTLV-I, which may explain why patient 5327 gave a tax+/pol− PCR pattern. We have seen this heterogenous pattern with the tax and pol primer detector systems previously in two ATL patients.17 In addition, patient 5048 consistently gave a stronger signal with tax than pol (Figs 1B and C) when the PCR products were assayed in the higher stringency liquid hybridization format. The presence of antibodies to the Tax and envelope proteins indicate that the provirus is transcriptionally active for these genes. It remains to be seen if there is some defect in the gag gene or its expression in these individuals; alternatively, this
phenomenon could be reflective of a normal expression pattern of the provirus. The phenomenon of having antibodies to regulatory proteins only may be reflective of the expression pattern of proviruses early after infection. It has been recently reported that some asymptomatic HIV patients who are HIV ELISA negative have antibodies against the regulatory NEF protein (Dr C. Baechot, World Health Organization Meeting, November 7-11, 1988, Paris). It is also quite possible that certain individuals do not, for one reason or another, develop an immune response to the gag proteins. It is highly unlikely, based on the compartmentalization of the Tax protein and the virus's life cycle, that these individuals did not have or do not have a productive infection.

It is formally possible that the PCR pol-negative patient is infected with a virus related to HTLV-I. If this is the case, however, it is not HTLV-II because this patient, as well as all the other patients with only anti-tax protein reactivity, was unreactive in an HTLV-II virion protein ELISA. This patient was also negative in an HTLV-II pol-specific amplification assay. His DNA was amplified with the HTLV-II-specific pol gene primers, SK58/59, and the amplified DNA was analyzed by liquid hybridization with SK60. We are currently attempting to follow up on these patients as well as the remaining anti-tax-positive patients.

The data reported herein double the confirmed exposure rate to HTLV-I within a cohort of white IDVAs and indicates that this phenomenon can occur in other at-risk populations. None of the ten sera were reactive in HTLV-I ELISAs prepared from viral proteins. These findings have important implications from both a public health perspective and for analyses of HTLV-I viral pathogenesis. We have developed a new ELISA that contains HTLV-I viral antigens and recombinant Tax protein. In preliminary trials this assay has produced a positive signal in a number of individuals in high-risk groups who were negative by the standard HTLV-I viral protein ELISA.

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