Purified Excreted-Secreted Antigens from *Trypanosoma cruzi*
Tryptomastigotes as Tools for Diagnosis of Chagas’ Disease

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There is currently no “gold standard” test for the diagnosis of late-stage Chagas’ disease. As a result, protection of the blood supply in areas where Chagas’ disease is endemic remains problematic. A panel of 709 serum samples from subjects with confirmed Chagas’ disease (n = 195), healthy controls (n = 400), and patients with other parasitic diseases (n = 114) was used to assess enzyme-linked immunosorbent assays (ELISAs) based on a concentrated extract of excretory-secretory antigens from either Brazil or Tulahuen strain *Trypanosoma cruzi* tryptomastigotes (total tryptomastigote excretory-secretory antigens [TESAs]). The total TESA-based assays had excellent overall sensitivity (100%) and specificity (>94%), except for cross-reactivity with *Leishmania*-infected sera. In an attempt to increase the specificity of the assay, immunoaffinity chromatography was used to purify the TESA proteins (TESA1A proteins). By Western blotting, a series of polypeptide bands with molecular masses ranging from 60 to 220 kDa were recognized by pooled sera positive for Chagas’ disease. An ELISA based on TESA1A proteins had a slightly lower sensitivity (98.6%) but an improved specificity (100%) compared to the sensitivity and specificity of the total TESA protein-based ELISAs. A 60-kDa polypeptide was identified as a major contributor to the cross-reactivity with *Leishmania*. These data suggest the need for field validation studies of TESA- and TESA1A-based assays in regions where Chagas’ disease is endemic.

*Trypanosoma cruzi* is the causative agent of Chagas’ disease, a major health problem in Central and South America (31). A large number of antigen preparations have been used for the serological diagnosis of this infection (1–5, 7, 8, 11, 12, 15, 18–20, 22, 27, 28). Of these preparations, tryptomastigote excretory-secretory antigens (TESAs) appear to provide good sensitivity and specificity as reagents for the diagnosis of Chagas’ disease. Major components of TESA proteins are transalilidases secreted into the culture supernatant (27, 29). Parasite transalilidases are implicated in the penetration of host cells as a result of the transfer of exogenous sialic acid to acceptor molecules located on the tryptomastigote surface. Transalilidases are highly immunogenic, and both the C-terminal and the N-terminal regions stimulate strong B-cell responses (9, 24). Transalilidase-like proteins are predominant antigens on the surfaces of bloodstream tryptomastigotes, metacyclic tryptomastigotes, and intracellular amastigotes (13).

Both enzyme-linked immunosorbent assay (ELISA) and immunoblot assays with TESA proteins have been reported for the diagnosis of Chagas’ disease. Immunoblots have typically been used in parasite diagnostics to rule out false-positive ELISA results (27). However, immunoblots are expensive, difficult to standardize, and time-consuming. In contrast, the ELISA format is inexpensive, simple to automate, and rapid (20). To date, the TESA proteins evaluated for use in the diagnosis of Chagas’ disease have been complex mixtures that cross-react with sera from subjects with other parasitic diseases, particularly leishmaniasis (28). We evaluated the performance of ELISAs based on concentrated TESA proteins or immunoaffinity chromatography-purified TESA (TESA1A) proteins for the diagnosis of latent Chagas’ disease. Both assays performed well in comparison with the performances of the serologic tests currently used in Central and South American blood banks. Of particular note, we identified a 60-kDa TESA protein that contributes significantly to the known cross-reactivity with *Leishmania* sera.

**MATERIALS AND METHODS**

**Serum samples.** A panel of 709 serum samples was used in this study. One hundred ninety-five samples were obtained from Venezuelan subjects with Chagas’ disease, as confirmed by a battery of three different serological tests, including immunofluorescence, indirect hemagglutination, and ELISA, at the National Chagas Immunodiagnosis Laboratory (NCIL; Maracay, Venezuela). Samples were considered positive if the results of two of the three assays were positive. Sera from 114 subjects with the following parasitic diseases were obtained from the Canadian National Reference Centre for Parasitology serum bank: leishmaniasis (n = 20), ascariasis (n = 6) fascioliasis (n = 8), malaria (n = 23), toxoplasmosis (n = 17), trichinosis (n = 11), filariasis (n = 8), cysticercosis (n = 8), and schistosomiasis (n = 13). These sera were not screened by the NCIL reference assays. Sera from 234 consecutive healthy Venezuelan blood donors negative by all three serological tests performed by NCIL and 166 healthy, nontraveling Canadians were also tested as controls.

**TESA proteins.** TESA proteins from two *T. cruzi* strains (strains Tulahuen and Brazil) were obtained from infected Vero cell supernatants, as described by Umezawa et al. (27), with slight modifications. Briefly, Vero cell monolayers at 65% confluence were infected with *T. cruzi* tryptomastigotes (1 × 10⁶ parasites/...
ml (175 cm²) and incubated at 37°C with 5% CO₂ for 4 days in Eagle’s minimum essential medium (Wisent, St. Bruno, Quebec, Canada) supplemented with 1 M HEPES (1%) and gentamicin reagent solution (0.1%) without fetal bovine serum and phenol red. After 4 days of incubation, the infected monolayers were washed twice with the medium and reincubated for 10 to 20 h at 37°C in 5% CO₂ in complete medium. The supernatants were then harvested, centrifuged at 2,800 x g for 15 min at 4°C, and filtered through a membrane (pore size, 0.22 µm; Millipore, Bradford, MA). The supernatant proteins were concentrated 32-fold (Ultra device; 30,000-molecular-weight cutoff; Amicon, Bradford, MA) and either used immediately or stored at −80°C. Total concentrated TESA proteins retained the high-molecular-mass (150- to 1,700-kDa) polypeptide bands, which correspond to the most immunogenic antigens (20). The protein content of the fetal bovine serum-free TESA was quantified by using a Micro BCA protein assay reagent kit (Pierce Co., Rockford, IL).

**Purification of chagasic antibodies.** Antibodies with specificity for T. cruzi were purified by following the procedure described by Curtis and Chase (10), with some modifications. A 10-ml pool of sera from subjects with serologically confirmed Chagas’ disease (NCIL assays) and no clinical or serological evidence of leishmaniasis was precipitated with 50% ammonium sulfate for 24 h and then centrifuged at 3,000 x g for 5 min. The resulting pellet was resuspended in 10 ml distilled water, and then 37% ammonium sulfate was added and the mixture was incubated for 24 h. After centrifugation at 3,000 x g for 5 min, the final pellet was resuspended in 10 ml of 0.02 M phosphate buffer (pH 8.0), and the residual ammonium sulfate was removed by dialysis (with 0.02 M phosphate buffer for 24 h). The final antibody concentration was determined by spectrophotometry by using the extinction coefficient (1.4) described previously (14).

Antibodies were purified by chromatography on a DEAE-cellulose column equilibrated with 0.02 M phosphate buffer (pH 8.0). Column elution fractions were monitored spectrophotometrically at 280 nm, and protein purity was determined by using sodium dodecyl sulfate (SDS)-polyacrylamide gel electrophoresis (PAGE). The fractions containing antibodies were pooled, and the final protein concentration was determined as reported by Bradford (6). Purified antibodies were dialyzed against 0.1 M NaHCO₃–0.5 M NaCl (pH 8.5) for 24 h and coupled to CNBr-activated Sepharose 4 Fast Flow, following the manufacturer’s instructions (Amersham Biosciences).

**Immunofinity purification of TESA proteins.** Aliquots of the total concentrated TESA (850 µg/ml) were mixed with the chagasic antibodies-Sepharose resin and incubated overnight in a nutator at 4°C. After the unbound proteins were washed with 0.1 M NaHCO₃–0.5 M NaCl (pH 8.5), the adsorbed antigens were eluted with 0.1 M glycine (pH 2.5). Fractions of 1 ml were collected in tubes containing 0.1 ml of 1 M Tris–HCl (pH 8.0) to neutralize the eluting buffer. Fractions were read at 280 nm, and the purity of the eluted proteins was assayed by SDS-PAGE, following by silver staining. Fractions containing the immunofinity chromatography-purified proteins (TESAα proteins) were dialyzed against phosphate-buffered saline (PBS; pH 7.4) for 24 h.

**TESA-based ELISAs.** Coating of 96-well polystyrene plates (Immulon 2; Thermo Labsystems, Franklin, MA) was accomplished by incubating either total TESAα (3 µg/ml) or TESAβ (3 µg/ml) for 6 h overnight at 4°C in 1 M sodium carbonate buffer (pH 9.6). The plates were washed four times with PBS (pH 7.4; 0.01 M phosphate buffer, 0.14 M NaCl) containing 0.05% Tween 20 (A&C, American Chemicals, Ltd., St-Laurent, Quebec, Canada) (PBST), blocked for 1 h at 37°C either with PBS containing 5% bovine serum albumin (Sigma, St. Louis, MO) and 0.1% Tween 20, or PBS containing 0.1% Tween 20, or PBS containing 5% skim milk (Parimal, Montreal, Quebec, Canada) and 0.1% Tween 20, or PBS containing 0.1% Tween 20 (in the case of TESAα). The sera were diluted 1:800 (total TESAα) or 1:200 (TESAβ) in the corresponding blocking solution (100 µl/well) and incubated for 1 h at 37°C. These dilutions were selected to permit optimal differentiation between positive and negative control sera, as tested by checkerboard titration (data not shown). Assays were completed with an optimal dilution of horseradish peroxidase-conjugated goat anti-human immunoglobulin G (IgG; 100 µl/well; Perkin-Elmer Life Science, Boston, MA) for 30 min at 37°C, four washes with PBST, and a final incubation with 3,3',5,5'-tetramethylbenzidine (100 µl/well; Serologicals Corporation, Massachusetts) for 10 min at room temperature. The reaction was stopped with 1 N H₂SO₄ (50 µl/well). The optical density (OD) was measured at 450 nm by using an automated ELISA reader (a Titertek Multiskan MCC/340 reader [Labsystems and Row Laboratories, Finland] or a TECAN ELISA reader). All experiments were performed in duplicate on different days, and pooled positive and negative controls for which the results were well established and included on each plate. The results were accepted only when the coefficient of variation within and among plates was ≤15%; otherwise, the samples were tested again. The results for healthy control sera were used to establish the cutoff values that yielded the optimal sensitivity and specificity results for each assay.

**RESULTS**

**Concentrated TESA fraction.** The total TESA was concentrated with an Amicon Ultra device that maintains the high-molecular-mass (150- to 1,700-kDa) immunogenic proteins (20). The final protein concentrations determined by using Micro BCA for TESA both before concentration (550 µg/ml) and after concentration (850 µg/ml) were similar for both strains. However, the OD values obtained from the strain Brazil- and the strain Tulahuen-based ELISAs before and after concentration were significantly different for a pool of positive sera by ANOVA (P values, <0.0001). Following concentration, the background activity for the negative control serum increased from ODs of 0.078 ± 0.021 and 0.059 ± 0.015 to ODs of 0.150 ± 0.018 and 0.130 ± 0.003 in the Tulahuen- and Brazil-based TESA ELISAs, respectively. In contrast, concentration of the TESAs increased the reactivity for the pooled positive sera 4.3-fold for the Tulahuen strain (OD range, 0.596 ± 0.033 to 2.162 ± 0.143) and 15-fold for the Brazil strain (OD range, 0.152 ± 0.013 to 2.237 ± 0.102).

**ELISA based on total TESA proteins.** The sensitivities of the ELISAs based on total TESA proteins were excellent for antigens from both strains (100%) at several cutoff values (OD cutoff values, 0.200 to 0.400) (Table 1). The assay based on the total TESA proteins derived from the Tulahuen strain showed a slightly higher specificity at the various cutoff values. However, when the results for all the negative control samples were combined for evaluation, these differences did not reach statistical significance. There were minor differences in the specificity estimates observed between the Tulahuen- and the Brazil-based assays for the individual control populations at various cutoff values (Table 2). However, the agreement between tests (kappa index) was excellent and in-

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**SDS-PAGE.** Protein purity was evaluated by SDS-PAGE carried out on 1.5-mm-thick slabs containing 7% polyacrylamide (for TESAα) or 11% polyacrylamide (for purified antibodies), as described by Laemmli (16). Electrophoresis was performed for 2 h in Tris-glycine electrode buffer (25 mM Tris, 192 mM glycine, 0.1% SDS, pH 8.3) at a constant voltage of 100 V. Coomassie blue R-250 (for purified antibodies) or silver staining (for TESAα) was used for protein visualization. Broad-range molecular weight markers (Promega, Madison, WI) and preestablished protein standards (Invitrogen, Carlsbad, CA) were included in each run.

**Western blot analysis.** Proteins separated by SDS-PAGE were transferred to nitrocellulose sheets (MFS, Pleasanton, CA), as described by Towbin et al. (26). Proteins were transferred by using a minitank electrophoret (Owl Scientific, Woburn, MA). Transfer was performed for 1 h at 4°C in 25 mM Tris, 192 mM glycine, and 20% (vol/vol) methanol (pH 8.3) at a current constant of 0.25 A. Membranes were blocked overnight at 4°C with PBS containing 5% skim milk (Parimal) and 0.1% Tween 20 and were incubated for 2 h with sera diluted 1:400 in the blocking solution. Following four 5-min washes with PBST, the antigenically reacting proteins were incubated for 2 h with the appropriate dilution of horseradish peroxidase-conjugated goat anti-human IgG diluted in PBS containing 5% skim milk and 0.1% Tween 20. The nitrocellulose filters were then washed four times with PBST, and the immune complexes were revealed by chemiluminescence (Super signal West Pico; Pierce).

**Statistical analyses.** Statistical analyses were performed by using SAS software (version 8.2; SAS, Cary, NC). Sensitivity and specificity were determined by examining all possible cutoff values. Analyses of variance (ANOVA; linear model) were used to examine the effect of parasite type on the optical density in each study group. Tukey’s post hoc test was used to evaluate differences between parasite antigens. Comparison among tests was determined by using the kappa index. P values of <0.05 were considered significant. The chi-square test was used to detect significant differences in specificity. The sensitivities and specificities of the tests were evaluated over a range of arbitrary OD cutoff values (OD value range, 0.2 to 0.4).

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TABLE 2. Mean, standard deviation, and range of OD values as well as specificity estimates for serum groups by ELISA based on TESAs from trypomastigote forms of *T. cruzi*.

<table>
<thead>
<tr>
<th>Patient group (n)</th>
<th>TESA source</th>
<th>Mean OD ± SD</th>
<th>OD range</th>
<th>Specificity (%) at an OD of:</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td></td>
<td></td>
<td></td>
<td>0.200</td>
</tr>
<tr>
<td>Canadian healthy patients (166)</td>
<td>B</td>
<td>0.069 ± 0.040</td>
<td>0.019–0.275</td>
<td>98.29</td>
</tr>
<tr>
<td></td>
<td>T</td>
<td>0.060 ± 0.032</td>
<td>0.018–0.181</td>
<td>100</td>
</tr>
<tr>
<td>Venezuelan healthy patients (234)</td>
<td>B</td>
<td>0.088 ± 0.060</td>
<td>0.031–0.820</td>
<td>97.44</td>
</tr>
<tr>
<td></td>
<td>T</td>
<td>0.087 ± 0.052</td>
<td>0.028–0.476</td>
<td>95.30</td>
</tr>
<tr>
<td>Patients with other parasitic diseases (114)</td>
<td>B</td>
<td>0.134 ± 0.200</td>
<td>0.027–2.077</td>
<td>94.02</td>
</tr>
<tr>
<td></td>
<td>T</td>
<td>0.107 ± 0.194</td>
<td>0.018–2.038</td>
<td>96.58</td>
</tr>
<tr>
<td>Chagas’ disease patients (195)</td>
<td>B</td>
<td>1.479 ± 0.506</td>
<td>0.450–2.628</td>
<td>98.29</td>
</tr>
<tr>
<td></td>
<td>T</td>
<td>1.558 ± 0.465</td>
<td>0.437–2.60</td>
<td>98.29</td>
</tr>
</tbody>
</table>

* a The mean ODs for patients with other parasitic diseases were as follows: for patients with ascariasis (n = 6), Brazil TESA assay, 0.145; Tulahuen TESA assay, 0.127; for patients with cysticeriosis (n = 8), Brazil TESA assay, 0.088; Tulahuen TESA assay, 0.064; for patients with filariasis (n = 8), Brazil TESA assay, 0.085; Tulahuen TESA assay, 0.070; for patients with trichinosis (n = 23), Brazil TESA assay, 0.133; Tulahuen TESA assay, 0.091; for patients with toxoplasmosis (n = 13), Brazil TESA assay, 0.073; Tulahuen TESA assay, 0.065.

* B, strain Brazil; T, strain Tulahuen.
Sera from healthy individuals and subjects with leishmaniasis did not react with the 170-, 120-, or 85-kDa polypeptide bands; but 3 of 10 of the serum samples from the leishmaniasis patients reacted with the 60-kDa polypeptide band. This observation suggests that this polypeptide is partially responsible for the known serologic cross-reactivity between these two parasitic diseases. Further evaluation with a larger number of serum samples from patients with leishmaniasis is required to confirm this result (Fig. 3).

ELISA based upon TESA$_{IA}$ proteins. We used a panel of 166 serum samples to perform a preliminary evaluation to compare the total TESA- and TESA$_{IA}$-based ELISAs (Tulahuen strain). Seventy-four of 75 serum samples from subjects with confirmed Chagas’ disease were reactive by the TESA$_{IA}$-based assay when an OD cutoff value of 0.300 was used (sensitivity, 98.7%; 95% confidence interval, 96 to 100%). None of the sera from healthy individuals from a region of endemicity ($n = 75$) or from patients with leishmaniasis ($n = 16$) were reactive in this assay (specificity, 100%). The mean OD values for chagasic sera, the negative control sera, and Leishmania-positive sera were $1.043 \pm 0.334$ (range, 0.298 to 1.735), $0.166 \pm 0.046$ (range, 0.086 to 0.284), and $0.145 \pm 0.048$ (range, 0.083 to 0.242), respectively. The mean OD for the Leishmania-positive sera was considerably lower in the optimized TESA$_{IA}$-based ELISA ($0.145 \pm 0.048$) compared to the mean values obtained in the total TESA-based assays with antigens from either the Tulahuen strain (0.253 ± 0.481) or the Brazil strain (0.300 ± 0.481). These differences did not reach statistical significance, however, due to the wide variability seen in the total TESA-based assays. Compared to the results of the optimized ELISAs based on total TESAs, the optimized TESA$_{IA}$ assay had a slightly lower sensitivity (98.7% versus 100%) but a higher specificity (100%) than the total TESA assays based on antigens from strains Brazil (97.8%) and Tulahuen (96.7%).

**DISCUSSION**

The use of TESA proteins for the diagnosis of Chagas’ disease has been reported, and their use has generally provided good results in terms of specificity and sensitivity (15, 18, 20, 27, 28). However, most studies have tested relatively small numbers of samples (18, 20, 28), and some have used TESA proteins from a single strain (27, 28). In the current study, we used a large panel of well-defined sera to evaluate ELISAs based on TESA proteins from two *T. cruzi* strains from different geographic regions (strains Brazil and Tulahuen). We demonstrated for the first time the utility of concentrating the most antigenic components of TESA, presumably by eliminating interfering low-molecular-weight peptides or proteins. In our hands, this simple manipulation dramatically increased the signal-to-noise ratio for the total TESA Brazil antigen in particular (~15-fold). We also show, for the first time, that immunoaffinity chromatography purification of TESA proteins can significantly enhance specificity with only a slight loss of sensitivity. Finally, our observations implicate a 60-kDa protein in the known serologic cross-reactivity that confounds the evaluation of patients and the screening of blood in regions where both Chagas’ disease and leishmaniasis are endemic.

We chose to test our concentrated total TESA ELISAs using samples obtained from a blood bank in a region where Chagas’ disease is endemic (i.e., in a real-life situation in which a candidate assay for Chagas’ disease might be used). Both the strain Brazil- and the strain Tulahuen-based assays had excellent sensitivities (100%) for the blood bank specimens confirmed to be positive and excellent specificities (99 to 100%) for samples from healthy control subjects from a region where Chagas’s disease is endemic and from a region where Chagas’s disease is not endemic. Furthermore, the tests maintained very reasonable specificities (94 to 99%) even when large numbers of samples from subjects with other parasitic diseases were tested. As has been reported previously (20, 28), sera from patients infected with *Leishmania* spp. were the most likely to cross-react in our TESA assays (2 of 20 presumed false-posit...
tive results for patients with cutaneous leishmaniasis in both the Brazil and the Tulahuen antigen-based assays). Moreover, our specificity estimate of ~90% for samples from leishmaniasis patients is likely to be an overestimate since no samples from patients with visceral leishmaniasis were included in our panel. Our data suggest that the concentrated total TESAs from both the Brazil and the Tulahuen strains include conserved epitopes that are highly immunogenic, which meets one of the principal criteria for Chagas’ disease tests suggested by the World Health Organization (32).

During the course of these studies, in addition to confirming the potential utility of the TESAs for the diagnosis of Chagas’ disease, we were also able to address several practical issues relevant to the potential use of these assays in the field. Large amounts of TESA proteins could be harvested with simple and inexpensive manipulations of relatively small volumes of trypanomastigote cultures (two 175-cm² Vero cell monolayers yielded ~5 ml of concentrated TESA proteins at ~850 µg/ml). Pooled total and concentrated TESA proteins were stable for at least 18 months when they were frozen as aliquots at ~80°C, with no loss of reactivity or increase in background “noise” (data not shown). The anticipated yield from only two infected 175-cm² flasks would permit the coating of approximately 420 ELISA plates and the performance of 16,800 tests. The 32-fold concentration of total TESA proteins that we achieved with a 30,000-molecular-weight cutoff filter resulted in marked increases in the signal-to-noise ratio, simplifying assay interpretation and the establishment of optimal cutoff values for specificity. It is likely that this procedure eliminates low-molecular-weight proteins-peptides that may interfere with the assay.

To our knowledge, the potential impact of immunoaffinity purification of TESAs has not been reported previously. In our hands, this step eliminated almost all of the complexity present in the total TESA protein mixtures, reducing the preparation from many bands to essentially five bands with molecular masses of 220, 170, 120, 85, and 60 kDa. The purification procedure consisted of a single step using Sepharose coupled to pooled antibodies from patients with Chagas’ disease. This approach resulted in a high yield of purified proteins (190 µg/ml from an initial 850-µg/ml concentration of TESAs) and had the advantage that the immunoaffinity column could be reused several times. Data for the TESAIA-based ELISA should be considered preliminary, however, because we performed only a small-scale purification that yielded enough material to test 166 of the 709 serum samples in our panel. A larger-scale purification of TESAIA and broader field studies of the TESAIA-based ELISA with different strains of T. cruzi are planned.

Many TESAs have recently been recognized to be transialidases. These enzymes may be virulence factors; and they participate in binding to noninfected epithelial, fibroblast, and muscle mammalian cell lines (9, 24). They are strong immunogens that elicit polyclonal antibody responses and are implicated in the induction of both inflammation and autoimmunity (21, 30). A monoclonal antibody (monoclonal antibody 39) that recognizes parasite transialidases detects polypeptide bands ranging in size from 85 to 170 kDa in TESA proteins (23). Several of these transialidase bands have molecular masses similar to those of the polypeptides present in our TESAIA preparation. Silber and colleagues (17, 25) used chromatography to purify T. cruzi proteins from total parasite extracts for the diagnosis of Chagas’ disease. They identified a 67-kDa lectin-like protein that binds to human erythrocyte membranes in a galactose-dependent fashion (25) and evaluated its potential as a diagnostic reagent (17). Although they reported a good sensitivity (98%) and a good specificity (98.11%) in an ELISA format, relatively small numbers of positive serum samples were tested, and only sera from healthy subjects were included as negative controls. Studies for the identification of the proteins present in our TESAIA preparation are under way.

To our knowledge, we are the first to implicate a 60-kDa protein in the serologic cross-reactivity between Leishmania-infected samples and samples from patients with Chagas’ disease. Cross-reactivity in an ELISA based on TESAs has previously been reported for subjects with visceral and cutaneous leishmaniases (20). In that study, Nakazawa and colleagues (20) implicated polypeptides of ≤150 kDa as cross-reactive but did not carry the characterization any further. Our TESAIA also contains a 60-kDa protein, yet no cross-reactivity was observed in the 16 Leishmania-infected samples tested, including the samples presumed to be false positive in the total TESA-based ELISA. The mean ODs for the Leishmania-positive sera were higher in the total TESA-based ELISAs than in the TESAIA-based ELISA, suggesting that immunoaffinity chromatography purification may eliminate the background “noise” that interferes with assay specificity. Alternately, the improved specificity of the TESAIA-based ELISA, despite the presence of a 60-kDa band in the TESAIA preparation by Western blotting, raises the possibility that two proteins of similar molecular masses are present in the total, concentrated TESA. It is also possible that presumably linear epitopes of a 60-kDa protein recognized by Leishmania-infected sera in the Western blot format are not present when TESAIA is used in the ELISA format. Studies are under way to determine whether or not removal of the reactive 60-kDa protein(s) from the total TESA preparation by molecular exclusion chromatography can achieve improved specificity while maintaining high sensitivity.

In summary, we have demonstrated that concentrated total TESA proteins are simple and inexpensive to produce, are stable at ~80°C for prolonged periods of time, and have an excellent sensitivity and an excellent specificity in an ELISA format. Using a simple protocol based on pooled chagasic antibodies, we have shown for the first time that immunoaffinity chromatography-purified TESA proteins can be used to significantly enhance the specificity of ELISA. These observations raise the possibility of testing by a sequential total TESA-based ELISA followed by a TESAIA-based ELISA as a simple and cost-effective strategy for screening of blood bank samples for Chagas’ disease.

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