- 1 Performance of commercially available serological screening tests for human T-cell
- 2 lymphotropic virus infection in Brazil
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- 4 Short title: Performance of screening tests for HTLV infection
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- 20
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- 23 **KEYWORDS** HTLV; Screening tests; Diagnostic reagent kits

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25	immunosorbent assay, particle agglutination or chemiluminescence assay kits. Due to antigen
26	matrix improvement entailing the use of new HTLV-antigens and changes in the format of
27	HTLV screening tests, as well as newly introduced CLIAs, a systematic evaluation of the
28	accuracy of currently available commercial tests is warranted. We aimed to assess the
29	performance of commercially available screening tests for HTLV diagnosis. A diagnostic
30	accuracy study was conducted on a panel of 397 plasma samples: 200 HTLV-negative, 170
31	HTLV-positive and 27 indeterminate under Western blotting analysis. WB-indeterminate
32	samples (i.e. those yielding no specific bands for HTLV-1 and/or HTLV-2) were assessed by
33	PCR and results were used to compare agreement among the commercially available ELISA
34	screening tests. For performance analysis, WB-indeterminate samples were excluded,
35	resulting in a final study panel of 370 samples. Three ELISA kits (Murex HTLV-1/2, anti-
36	HTLV-1/2 SYM Solution and Gold ELISA HTLV-1/2) and one CLIA kit (Architect r-
37	HTLV-1/2) were evaluated. All screening tests demonstrated 100% sensitivity. Concerning
38	the HTLV-negative samples, SYM Solution and Gold ELISA kits had specificity values
39	>99.5%, while the Architect r-HTLV-1/2 test presented 98.1% specificity, followed by
40	Murex (92.0%). Regarding the 27 samples with WB-indeterminate results, after PCR
41	confirmation, all ELISA kits showed 100% sensitivity, but low specificity. Accuracy findings
42	were corroborated by Cohen's Kappa, which evidenced slight and fair agreement between
43	PCR analysis and ELISA tests for HTLV diagnosis. Based on the data, we believe that all
44	evaluated tests can be safely used for HTLV-infection screening.
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ABSTRACT Serological screening for HTLV-1 is usually performed using enzyme-linked

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Human T-cell lymphotropic virus type 1 (HTLV-1) and type 2 (HTLV-2) were identified in
1980 and 1982, respectively (1, 2). Subsequently, HTLV-3 and HTLV-4 were discovered in
2005 (3, 4). It has been estimated that at least 10 million people harbor HTLV-1 worldwide
(5). Large foci of this infection exist in Japan, Africa, the Caribbean Islands, Melanesia,
Australia, the Mashhad area of northeastern Iran and South America (5–7). HTLV-1 is
associated with, or causes, a broad range of inflammatory conditions and a severe
proliferative disease (5, 8–14).

HTLV-2 infection is endemic in native Amerindian populations in both North and South America, certain tribes of Pygmies in Africa and in intravenous drug users (IDUs) in urban areas around the world (15, 16). In contrast to HTLV-1, this type rarely is associated with neurological or lymphoproliferative disorders (17). HTLV-3 and HTLV-4 are restricted to Western Africa and have not yet been associated with any diseases (3, 4).

Brazil, a country of 200 million inhabitants, has a population of 800,000 who potentially harbor HTLV-1, representing one of the largest endemic areas for the virus and its associated diseases anywhere in the world (5). The virus is disseminated throughout the country, with higher rates found in the Northeast and Northern regions compared with the South and Southeast (18, 19). HTLV-2 is present mainly in the North, among indigenous populations and in IDUs in urban centers (17).

Achieving an accurate diagnosis of HTLV infection is a complex task. Serological screening for HTLV-1 is usually performed using enzyme-linked immunosorbent assay (ELISA), particle agglutination testing or chemiluminescence assay (CLIA) kits. The Brazilian Ministry of Health recommends the use of ELISA or particle agglutination tests as a screening protocol. Western blotting (WB) or immunoblot is used for confirmation, and polymerase chain reaction (PCR) is employed in the case of inconclusive confirmatory test

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results (20). Among the screening options, ELISA is used most extensively due to an elevated level of automation, simplicity and low cost. ELISA performance depends on antigen composition and assay format (21–24). Tests providing low accuracy present a public health problem, as false-positive results can have a negative impact, not only economically due to the need for confirmation by WB, but also on individuals' quality of life.

In light of this scenario, we endeavored to conduct a systematic evaluation of the commercial screening test kits for HTLV diagnosis. Statistical tools were used to obtain a robust assessment of the performance of each molecule by determining the following diagnostic test parameters: sensitivity (probability of test being positive in the presence of infection) and specificity (probability of test being negative in the absence of infection).

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82 MATERIAL AND METHODS

Ethical considerations. The present research protocol was approved by the Institutional Research Board (IRB) of the Bahiana School of Medicine and Public Health (EBMSP) in Salvador (protocol no. 464.286). All procedures were performed in accordance with the principles established in the Declaration of Helsinki and its subsequent revisions.

87 Sample selection. The present diagnostic accuracy study was carried out between February 2015 and December 2017 using anonymous plasma samples obtained from the 88 biorepository of the Integrated and Multidisciplinary HTLV Center (CHTLV) at EBMSP. 89 CHTLV is an outpatient clinic, open to the public, that provides inter-disciplinary care and 90 91 services, including general medical treatment, laboratory diagnosis, psychological 92 counseling and physical therapy. All included plasma samples had been previously screened for antibodies against HTLV-1/2 using an enzyme-linked immunosorbent assay (Ortho® 93 HTLV-1/HTLV-2 Ab-Capture ELISA Test Systems, Ortho-Clinical Diagnostic, Raritan, 94

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USA), and reactive samples were retested by Western Blot (HTLV Blot 2.4, Genelabs 95 Diagnostics[®], Singapore). Test results were interpreted according to the stringent criteria 96 indicated by the manufacturer and in accordance with the guidelines established by the 97 Brazilian Ministry of Health (20). 98

The panel consisted of 397 samples: 200 HTLV-negative, 170 HTLV-positive (122 99 100 HTLV-1, 31 HTLV-2, 5 HTLV-1+HTLV-2, and 12 HTLV), and 27 WB-indeterminate. Briefly, HTLV-negative samples were defined as those lacking reactivity to HTLV-specific 101 102 proteins; HTLV-1-positive samples were defined as reactive to GAG (p19 with or without 103 p24) and two ENV (GD21 and rgp46-I); HTLV-2-positive samples were reactive to GAG (p24 with or without p19) and two ENV (GD21 and rgp46-II); HTLV seropositive samples 104 105 were reactive to GAG (p19 and p24) and ENV (GD21); samples were considered indeterminate when no HTLV specific bands were detected, i.e. the criteria for HTLV-I, 106 HTLV-II or HTLV were not satisfied. Indeterminate samples were assessed by PCR analysis 107 108 and the obtained results were used to compare agreement with ELISA screening test results. For performance analysis, the WB-indeterminate samples were excluded, forming a final 109 study panel of 370 samples (Fig. 1). 110

Alternatively, 217 plasma samples (112 positive, 105 negative) were also assessed by 111 112 chemiluminescence assay - CLIA (Architect rHTLV-1/2, Abbott Diagnostics Division, Wiesbaden, Germany). 113

Immunoassays. Three HTLV1/2-specific enzyme immunoassay kits, 114 all 115 commercially available in Brazil, were employed in this study: Murex HTLV-1/2 (DiaSorin S.p.A., Dartford, UK), anti-HTLV-1/2 Sym Solution (Symbiosis Diagnostica LTDA, Leme, 116 117 Brazil) and Gold ELISA HTLV-1/2 (Rem Indústria e Comércio LTDA, São Paulo, Brazil). 118 Cut-off values, as well as gray zones, were calculated for each test as follows: by adding 0.2 119 to the mean of the negative control replicates for Murex HTLV-1/2; adding 0.18 to the mean 120 of the negative control replicates for Anti-HTLV-1/2 Sym Solution; by adding 0.25 to the mean of the negative control replicate for Gold ELISA HTLV-1/2. For data normalization, 121 122 all results were expressed by plotting values in an indexed format, calculated as the ratio 123 between a given sample's optical density (OD) and the cut-off OD values respective to each 124 assay. Under this index, referred to as a reactivity index (RI), all results <1.00 were considered negative. When a sample's RI value was $1.0 \pm 10\%$, the result was considered as 125 126 indeterminate (i.e. in the grey zone), and these samples were deemed inconclusive.

127 HTLV-1/2 molecular detection. Peripheral blood mononuclear cells (PBMC) from 27 patients with WB-indeterminate results were obtained from EDTA blood samples under 128 129 density gradient centrifugation; DNA was extracted using a spin column kit (Qiagen, Hilden, Germany). DNA samples were submitted to nested-PCR using the HTLV-1 long terminal 130 repeat (LTR) 5' region primers as described previously (25), outer primers BSQF6/BSDR3 131 132 and inner primers BSQF2/BSDR4, to amplify a 672-bp fragment in the HTLV-2 LTR region (26). All amplified products were submitted to electrophoresis on a 1% agarose gel with 133 134 Syber Safe DNA (Invitrogen).

Statistical analysis. Data were encoded and analyzed using scatterplot computer 135 136 graphic software (Prism version 7; GraphPad, San Diego, CA). Descriptive statistics are presented as geometric means \pm standard deviation. To test data normality, the Shapiro-Wilk 137 138 test, followed by Student's t-test, were used. When assumed homogeneity was not confirmed, 139 Wilcoxon's signed rank test was used. All analyses were two-tailed, and p-values under 5% 140 were considered significant (p < 0.05). Enzyme immunoassay test performance was 141 computed using a dichotomous approach and compared in terms of sensitivity, specificity, 142 accuracy, likelihood ratio (LR) and diagnostic odds ratio (DOR). Additionally, receiver

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143	operating characteristic (ROC) curves were constructed and the areas under these curves
144	were used as a global measure of test performance. Confidence intervals (CI) were employed
145	at a confidence level of 95%. The strength of agreement between screening commercial tests
146	and PCR results was assessed by Cohen's Kappa coefficient (κ) (27), which accounts for
147	agreement occurring only by chance beyond simple percentage agreement calculations. $\boldsymbol{\kappa}$
148	values are interpreted as poor ($\kappa \le 0$), slight ($0 \le \kappa \le 0.20$), fair ($0.21 \le \kappa \le 0.40$), moderate
149	$(0.41 < \kappa \le 0.60)$, substantial $(0.61 < \kappa \le 0.80)$ and almost perfect agreement $(0.81 < \kappa \le 1.0)$.
150	A flowchart (Fig. 1) have been provided the Standards for Reporting of Diagnostic Accuracy
151	Studies (STARD) guidelines (28).

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153 **RESULTS**

Assay performance. Using plasma from 170 HTLV-positive individuals, ELISA and
CLIA performance were assessed, as shown in Fig. 2. The area under the curve (AUC) values
were >99%, demonstrating excellent overall diagnostic accuracy for all kits tested. RI values
for HTLV-1/2-positive samples were variable, ranging from 14.2 for SYM Solution, 14.5 for
Gold ELISA to 16.8 for Murex. In addition, Architect r-HTLV-1/2 yielded the highest RI
value (>90).

As all kits test demonstrated 100% sensitivity, no statistically significant differences were detected. Regarding the HTLV-1/2-negative samples, SYM Solution and Gold ELISA presented specificity values >99%. Architect r-HTLV-1/2 showed a specificity of 98.1%, followed by Murex at 92.0%. Differences in specificity and RI were not statistically significant between the SYM Solution and Gold ELISA kits. With respect to HTLV-negative samples, the maximum RI value was obtained using Murex (RI = 0.30) (Fig. 2). Considering

166 RI values of 1.0 ± 0.10 as inconclusive, i.e. falling in the gray zone, we verified that truly 167 positive HTLV-1/2 samples were conclusively diagnosed by the Gold ELISA, Murex, SYM Solution and Architect r-HTLV-1/2 tests. As regards the HTLV-negative samples, one fell 168 in the gray zone using the Gold ELISA test. With respect to diagnostic accuracy, Gold 169 ELISA, SYM Solution, and Architect rHTLV-1/2 tests demonstrated the highest accuracy 170 171 (>99.1%), while Murex presented the lowest result (95.6%). DOR scores, based on likelihood ratios, were 524,000 for Architect rHTLV-1/2, 338,200 for Gold ELISA, 168,254 for SYM 172 173 Solution and 19,552 for Murex HTLV-1/2. Among the ELISA kits evaluated, Gold ELISA 174 offered the best performance, as evidenced by ROC analysis and, notably, the exceptionally 175 high diagnostic odds ratio produced by this test (Fig. 2). No significant differences in RI signal were observed with regard to the different types of seroreactivity (HTLV-1 vs. HTLV-176 2 vs. HTLV-1/2 and HTLV). 177

Assay agreement. Analysis of the diagnostic accuracy of the three commercial 178 179 ELISAs with respect to 27 WB-indeterminate samples, considering PCR amplification as a gold standard for HTLV diagnosis, revealed that eight samples were negative (29.6%) and 180 19 were positive (70.4%) for HTLV-1 (Fig. 3), with all ELISA tests yielding 100% 181 182 sensitivity. Conversely, all three assays presented specificity inferior to 25%, with Gold 183 ELISA offering just 12.5% specificity. Despite this very low accuracy, both Murex and SYM Solution kits offered higher accuracy than Gold ELISA. Slight and fair agreement (Cohen's 184 Kappa < 0.40) between PCR analysis and the ELISA screening tests was detected with regard 185 186 to diagnosing HTLV infection. Table 1 details the 27 HTLV-indeterminate profiles that 187 allowed for the identification of distinct patterns. No HGIP (29) or N (30) patterns were 188 observed.

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191 The present study found a high diagnostic value for each of the four different evaluated commercially available HTLV screening tests used to detect anti-HTLV antibodies 192 in Brazil. In fact, AUC values greater than 99% demonstrates convincing evidence of the 193 optimal discriminative power of these kits regarding HTLV-positive and HTLV-negative 194 195 samples. Gold ELISA and Architect rHTLV-1/2 both presented AUC values of 100%. Furthermore, the Murex, SYM Solution and Architect rHTLV-1/2 assays did not show 196 197 inconclusive results (grey zone) in HTLV antibody screening procedures. Gold ELISA 198 yielded low number of inconclusive results, as only one out of 170 HTLV-positive samples tested using this kit produced an RI value that fell in the grey zone. 199

All tests displayed 100% sensitivity in diagnosing HTLV-positive samples. RI values were higher than 14 for the ELISA tests and above 90 for the Architect rHTLV-1/2 kit, which corroborates previous reports (31). Regarding the ELISA tests, the highest RI value was achieved by Murex, with statistically significant differences seen compared to Gold ELISA and SYM Solution.

Due to the high number of misdiagnosed samples (4.3%) under the Murex test, its 205 206 accuracy was significantly lower compared to the other kits. Gold ELISA, SYM Solution and 207 Architect rHTLV-1/2 were all found to be 99% accurate, suggesting that these kits can be safely employed for HTLV infection screening. Although the Murex test is less accurate, it 208 nonetheless returned values above 95%, indicating suitability in the diagnosis of HTLV 209 210 infection; however, the proportion of samples requiring WB confirmation was greater, which 211 increases the cost of performing diagnosis. In fact, 8% of the HTLV-negative samples 212 assayed with Murex yielded false-positive results, with a specificity of 92%. It is interesting 213 to note that this test's performance has improved over time, as studies performed in 2007 and 214 2009 described its sensitivity and specificity at 98.2% and 42.6%, respectively (32, 33). 215 Another study conducted in Argentina showed that Murex was 97.2% sensitive and 99.7% specific (34). More recently, other studies have reported high values of specificity, such as 216 those evaluating HIV/HTLV co-infected individuals (99.0%) (31) and blood donors (97.2%) 217 218 (34). With respect to HTLV-negative samples, the Murex test returned the highest RI value. 219 The observed differences in RI values could arise from variability in antigenic composition. While all tests correctly diagnosed positive samples, it is possible that the antigenic matrix 220 221 employed in the solid phase of the Murex kit recognized no specific anti-HTLV antibodies, 222 which led to false-positive results or cross-reactions.

223 Of note, the sensitivity, specificity, and accuracy values associated with diagnostic 224 tests are unsatisfactory in terms of influencing clinical decisions (35). A diagnostic test can 225 only be considered valid if the results produced modify the probability of disease occurrence. Likelihood Ratio (LR) measurements can be helpful in describing a test's discriminatory 226 227 power and determining the possibility of a particular result occurring among infected individuals, as opposed to the probability of the same result being obtained among healthy 228 individuals (36). In our study, Gold ELISA had a positive LR of 201, indicating that an 229 HTLV-infected person is approximately 201 times more likely to be diagnosed with this 230 231 infection if evaluated with this kit. The lowest positive LR value was observed with the Murex test (12.6), indicating a low probability for an HTLV-infected person to be accurately 232 diagnosed. Conversely, a study performed in 2008 found a positive LR of 326.5 for Murex 233 234 (34). HTLV-negative samples returned LR values lower than 0.001 under all of the evaluated 235 tests. There is a consensus that positive LRs above 10 and negative LRs below 0.1 contribute 236 substantially to diagnosis (36). DOR, calculated as the ratio between positive and negative 237 LR values, is considered a global performance parameter that summarizes the diagnostic test

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238 accuracy. DOR values describe the probability of receiving a positive result for a person with 239 infection, as opposed to someone who is uninfected (35). The DOR for Architect rHTLV-1/2 (524,000) was the highest among the screening tests evaluated, followed by Gold ELISA 240 (338,200), SYM Solution (168,254) and Murex (19,552). These findings suggest that 241 Architect rHTLV-1/2 and Gold ELISA offer superior performance to SYM Solution and 242 243 Murex. LR and DOR determination are relevant and stable tools, since these parameters remain independent of the prevalence of disease (37). The HTLV-1 and HTLV-2-244 245 seroindeterminate WB patterns observed herein were similar to those reported by other 246 studies. However, no HGIP or N patterns were identified.

247 It is important to note that, concerning the Architect rHTLV-1/2 test, our findings are 248 in agreement with those reported by other studies. In fact, identical values of sensitivity (100%) and specificity (>99%) have been described in both samples from blood donors and 249 hospitalized patients (38). Similar results were demonstrated by Malm et al. (39) (Sen 100%; 250 251 Spe 99.8%), as well as by Qiu et al. (40) (Sen 100%; Spe 99.98%) in general populations of the USA, Japan and Nicaragua. Although the present study was unable to assess other 252 253 screening tests, the literature indicates the high performance of both the Elecsys HTLV-I/II and Abbott Prism HTLV-I/HTLV-II kits (Sen 100%; Spe > 99%) in samples from both blood 254 255 donors and other obtained from a routine diagnostic service (41). The DiaSorin LIAISON® XL recHTLV-I/II kit was also evaluated elsewhere, with high sensitivity and specificity 256 values reported, similarly to the Architect rHTLV-1/2 test (42-44). 257

258 The results presented herein indicate that all evaluated kits can safely be used for HTLV-259 infection screening. However, it is important to note that the high sensitivity offered by these 260 kits may lead to false-positive results, which could increase cost as a result of WB 261 confirmation requirements. From the perspective of large diagnostic centers and blood banks,

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262	proper screening method selection can substantially reduce costs associated with
263	confirmatory testing. In an effort to reduce costs and assure correct diagnosis, a new
264	diagnostic protocol for HTLV-infection diagnosis was proposed by Costa et al. (45), who
265	suggested the use of two ELISA tests for screening purposes, followed by real-time PCR. In
266	this case, WB confirmation would only be indicated in cases of negative PCR results. Herein,
267	when the 27 WB-indeterminate samples were analyzed by PCR, all HTLV-1 positive samples
268	demonstrated agreement with results from each of the three ELISA tests evaluated. On the
269	other hand, overall agreement was slight or fair due to the high number of false-positive
270	results obtained using ELISA. Moreover, it has been demonstrated that the INNO-LIA HTLV
271	I/II Ab serological confirmatory assay for HTLV yielded results for most of the samples
272	considered indeterminate or positive, but untypeable, in WB assays (31, 46). These data
273	suggest the costs associated with HTLV-infection diagnosis could be lowered by using
274	molecular biology-based methodologies, or INNO-LIA HTLV, as a confirmatory assay in
275	place of WB. In the context of low-income countries, such as those in Africa and Latin
276	America, we suggest that CLIA represents a suitable screening strategy for blood banks due
277	to the high DOR values found herein. However, in countries lacking the necessary
278	infrastructure, the use of an ELISA offering a high DOR value, e.g. Gold Elisa, seems to be
279	a satisfactory alternative.

Despite the scarcity of studies evaluating the diagnostic performance of screening tests in 280 diagnosing HTLV-infection by employing LR, DOR and AUC as performance parameters, 281 we evaluated three ELISA tests and one CLIA used for HTLV-infection screening. Based on 282 283 the present findings, we conclude that all of the 3rd generation commercially available kits 284 employed herein presented high sensitivity and specificity values compared to previous

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studies. Among the ELISA tests evaluated, the Gold HTLV-1/2 kit offered the best 285 286 performance parameters, while the ARCHITECT rHTLV-1/2 demonstrated the highest performance of all the assays considered. High sensitivity values produced by screening tests 287 288 could lead to high proportions of false-positive results. Thus, we reinforce our previous suggestion and urge the consideration of a new protocol employing molecular biology or line 289 290 immune assay (INNO-LIA HTLV) techniques as a first choice for confirmatory testing in place of WB. 291

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ACKNOWLEDGEMENTS 293

294 We thank Andris K. Walter for providing English language revision and manuscript 295 copyediting assistance. This work was supported by Conselho Nacional de Desenvolvimento Científico e Tecnológico - CNPq (473667/2012-6 and 311054/2014-5), Fundação de Amparo 296 à Pesquisa do Estado da Bahia - FAPESB (2574/2013), and Fundação Nacional para o 297 Desenvolvimento do Ensino Superior - FUNDADESP (9600113). 298

299

300 Competing interests: The authors have declared that no competing interests exist. 302 1. Poiesz BJ, Ruscetti FW, Gazdar AF, Bunn PA, Minna JD, Gallo RC. 1980. Detection and isolation of type C retrovirus particles from fresh and cultured 303 lymphocytes of a patient with cutaneous T-cell lymphoma. Proc Natl Acad Sci USA 304 77:7415-7419. https://www.ncbi.nlm.nih.gov/pmc/articles/PMC350514. 305 306 2. Kalyanaraman VS, Sarngadharan MG, Robert-Guroff M, Miyoshi I, Golde D, Gallo RC. 1982. A new subtype of human T-cell leukemia virus (HTLV-II) associated with 307 308 a T-cell variant of hairy cell leukemia. Science 218:571-573. https://doi.org/ 309 10.1126/science.6981847. 3. Calattini S, Chevalier SA, Duprez R, Bassot S, Froment A, Mahieux R, Gessain A. 310 311 2005. Discovery of a new human T-cell lymphotropic virus (HTLV-3) in Central Africa. Retrovirology 2:30. https://doi.org/10.1186/1742-4690-2-30. 312 4. Wolfe ND, Heneine W, Carr JK, Garcia AD, Shanmugam V, Tamoufe U, Torimiro 313 314 JN, Prosser AT, Lebreton M, Mpoudi-Ngole E, McCutchan FE, Birx DL, Folks TM, Burke DS, Switzer WM. 2005. Emergence of unique primate T-lymphotropic viruses 315 among central African bushmeat hunters. Proc Natl Acad Sci USA 102:7994-7999. 316 317 https://doi.org/10.1073/pnas.0501734102. 5. Gessain A, Cassar O. 2012. Epidemiological Aspects and World Distribution of 318 HTLV-1 Infection. Front Microbiol 3:388. 319 https://doi.org/10.3389/fmicb.2012.00388. 320 321 6. Rafatpanah H, Hedayati-Moghaddam MR, Fathimoghadam F, Bidkhori HR, 322 Shamsian SK, Ahmadi S, Sohgandi L, Azarpazhooh MR, Rezaee SA, Farid R, 323 Bazarbachi A. 2011. High prevalence of HTLV-I infection in Mashhad, Northeast Iran: a population-based seroepidemiology survey. J Clin Virol 52:172-176. 324

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325		https://doi.org/10.1016/j.jcv.2011.07.004.
326	7.	Einsiedel L, Purcell D, Schinke S, Haynes K, Taylor GP, Martin F. 2018. Highlights
327		from the HTLV-1 symposium at the 2017 Australasian HIV and AIDS Conference
328		held jointly with the 2017 Australasian Sexual Health Conference, November 2017,
329		Canberra, Australia. J virus Erad 4:48-50.
330		https://www.ncbi.nlm.nih.gov/pmc/articles/PMC5851185.
331	8.	Yoshida M, Miyoshi I, Hinuma Y. 1982. Isolation and characterization of retrovirus
332		from cell lines of human adult T-cell leukemia and its implication in the disease.
333		Proc Natl Acad Sci USA 79:2031-2035. https://doi.org/10.1073/pnas.79.6.2031.
334	9.	Gessain A, Vernant JC, Maurs L, Barin F, Gout O, Calender A, De Thé G. 1985.
335		Antibodies to human T-lymphotropic virus type-I inpatients with tropical spastic
336		paraparesis. Lancet 326:407-410. https://doi.org/10.1016/S0140-6736(85)92734-5.
337	10.	Osame M, Usuku K, Izumo S, Ijichi N, Amitani H, Igata A, Matsumoto M, Tara M.
338		1986. HTLV-I associated myelopathy, a new clinical entity. Lancet 1:1031-1032.
339		https://doi.org/10.1016/S0140-6736(86)91298-5.
340	11.	LaGrenade L, Hanchard B, Fletcher V, Cranston B, Blattner W. 1990. Infective
341		dermatitis of Jamaican children: a marker for HTLV-I infection. Lancet 336:1345-
342		1347. https://doi.org/10.1016/0140-6736(90)92896-P.
343	12.	Mochizuki M, Watanabe T, Yamaguchi K, Yoshimura K, Nakashima S, Shirao M,
344		Araki S, Takatsuki K, Mori S, Miyata N. 1992. Uveitis associated with human T-cell
345		lymphotropic virus type I. Am J Ophthalmol 114:123-129.
346		https://doi.org/10.1016/S0002-9394(14)73974-1.
347	13.	Castro-Lima Vargens C, Grassi MFR, Boa-Sorte N, Rathsam-Pinheiro RH, Olavarria
348		VN, de Almeida Kruschewsky R, Galvão-Castro B. 2011. Keratoconjunctivitis sicca

349		of human T cell lymphotropic virus type 1 (HTLV-1) infected individuals is
350		associated with high levels of HTLV-1 proviral load. J Clin Virol 52:177-180.
351		https://doi.org/10.1016/j.jcv.2011.07.016.
352	14.	Einsiedel L, Pham H, Wilson K, Walley R, Turpin J, Bangham C, Gessain A,
353		Woodman RJ. 2018. Human T-Lymphotropic Virus type 1c subtype proviral loads,
354		chronic lung disease and survival in a prospective cohort of Indigenous Australians.
355		PLoS Negl Trop Dis 12:e0006281. https://doi.org/10.1371/journal.pntd.0006281.
356	15.	Araujo A, Hall WW. 2004. Human T-lymphotropic virus type II and neurological
357		disease. Ann Neurol 56:10-19. https://doi.org/10.1002/ana.20126.
358	16.	Mauclère P, Afonso PV, Meertens L, Plancoulaine S, Calattini S, Froment A, Van
359		Beveren M, de Thé G, Quintana-Murci L, Mahieux R, Gessain A. 2011. HTLV-2B
360		strains, similar to those found in several Amerindian tribes, are endemic in central
361		African Bakola Pygmies. J Infect Dis 203:1316-1323.
362		https://doi.org/10.1093/infdis/jir031.
363	17.	Paiva A, Casseb J. 2015. Origin and prevalence of human T-lymphotropic virus type
364		1 (HTLV-1) and type 2 (HTLV-2) among indigenous populations in the Americas.
365		Rev Inst Med Trop Sao Paulo 57:1-13. https://doi.org/10.1590/S0036-
366		46652015000100001.
367	18.	Galvão-Castro B, Loures L, Rodriques LG, Sereno A, Ferreira Júnior OC, Franco
368		LG, Muller M, Sampaio DA, Santana A, Passos LM, Proietti F. 1997. Distribution of
369		human T-lymphotropic virus type I among blood donors: a nationwide Brazilian
370		study. Transfusion 37:242-243. https://doi.org/10.1046/j.1537-
371		2995.1997.37297203532.x.
372	19.	Catalan-Soares B, Carneiro-Proietti AB de F, Proietti FA, Interdisciplinary HTLV

373		Research Group. 2005. Heterogeneous geographic distribution of human T-cell
374		lymphotropic viruses I and II (HTLV-I/II): serological screening prevalence rates in
375		blood donors from large urban areas in Brazil. Cad Saude Publica 21:926-931.
376		https://doi.org/10.1590/S0102-311X2005000300027.
377	20.	Brasil. Ministério da Saúde. Secretaria de Vigilância em Saúde. Programa Nacional
378		de DST e Aids. 2003. Guia do manejo clínico do HTLV / Ministério da Saúde,
379		Secretaria de Vigilância em Saúde, Programa Nacional de DST e Aids, p. 41. In
380		Ministério da Saúde, Brasília. Available from:
381		http://bvsms.saude.gov.br/bvs/publicacoes/guia_de_manejo_clinico_do_paciente_co
382		m_HTLV.pdf.
383	21.	Hartley TM, Malone GE, Khabbaz RF, Lal RB, Kaplan JE. 1991. Evaluation of a
384		recombinant human T-cell lymphotropic virus type I (HTLV-I) p21E antibody
385		detection enzyme immunoassay as a supplementary test in HTLV-I/II antibody
386		testing algorithms. J Clin Microbiol 29:1125-1127.
387		https://jcm.asm.org/content/29/6/1125.long.
388	22.	Soriano V, Gutiérrez M, Tuset C, Martínez-Zapico R, Calderón E, González-Lahoz
389		J. 1995. Avoiding false-negative results for HTLV-II using new serological assays.
390		Am J Med 98:103. https://doi.org/10.1016/S0002-9343(99)80095-3.
391	23.	Andersson S, Thorstensson R, Ramirez KG, Krook A, von Sydow M, Dias F,
392		Biberfeld G. 1999. Comparative evaluation of 14 immunoassays for detection of
393		antibodies to the human T-lymphotropic virus types I and II using panels of sera
394		from Sweden and West Africa. Transfusion 39:845-851.
395		https://doi.org/10.1046/j.1537-2995.1999.39080845.x.
396	24.	Caterino-de-Araujo A. 2009. Best screening assays for the diagnosis of human T-cell

397		lymphotropic virus types 1 and 2 in South America. J Virol Methods 156:150-151.
398		https://doi.org/10.1016/j.jviromet.2008.10.003.
399	25.	Mboudjeka I, Zekeng L, Yamashita M, Takehisa J, Ido E, Miura T, Ohkura S, Ikeda
400		M, Kaptue L, Hayami M. 1997. Prevalence and phylogenetic analysis of HTLV-I
401		isolates in Cameroon, including those of the Baka Pygmy. Jpn J Cancer Res 88:619-
402		624. https://doi.org/10.1111/j.1349-7006.1997.tb00427.x.
403	26.	Alcantara LCJ, Shindo N, Van Dooren S, Salemi M, Costa MCR, Kashima S, Covas
404		DT, Vandamme A-M, Galvão-Castro B. 2003. Brazilian HTLV type 2a strains from
405		intravenous drug users (IDUs) appear to have originated from two sources: Brazilian
406		Amerindians and European/North American IDUs. AIDS Res Hum Retroviruses
407		19:519-523. https://doi.org/10.1089/088922203766774577.
408	27.	Landis JR, Koch GG. 1977. The measurement of observer agreement for categorical
409		data. Biometrics 33:159-174. https://www.ncbi.nlm.nih.gov/pubmed/843571.
410	28.	Bossuyt PM, Reitsma JB, Bruns DE, Gatsonis CA, Glasziou PP, Irwig L, Lijmer JG,
411		Moher D, Rennie D, de Vet HCW, Kressel HY, Rifai N, Golub RM, Altman DG,
412		Hooft L, Korevaar DA, Cohen JF. 2015. STARD 2015: an updated list of essential
413		items for reporting diagnostic accuracy studies. BMJ h5527.
414		https://doi.org/10.1136/bmj.h5527.
415	29.	Rouet F, Meertens L, Courouble G, Herrmann-Storck C, Pabingui R, Chancerel B,
416		Abid A, Strobel M, Mauclere P, Gessain A. 2001. Serological, epidemiological, and
417		molecular differences between human T-cell lymphotropic virus Type 1 (HTLV-1)-
418		seropositive healthy carriers and persons with HTLV-I Gag indeterminate Western
419		blot patterns from the Caribbean. J Clin Microbiol 39:1247-1253.
420		https://doi.org/0.1128/JCM.39.4.1247-1253.2001.

18

421	30.	Filippone C, Bassot S, Betsem E, Tortevoye P, Guillotte M, Mercereau-Puijalon O,
422		Plancoulaine S, Calattini S, Gessain A. 2012. A new and frequent human T-cell
423		leukemia virus indeterminate Western blot pattern: epidemiological determinants and
424		PCR results in central African inhabitants. J Clin Microbiol 50:1663-1672.
425		https://doi.org/10.1128/JCM.06540-11.
426	31.	Campos KR, Gonçalves MG, Costa NA, Caterino-de-Araujo A. 2017. Comparative
427		performances of serologic and molecular assays for detecting human T lymphotropic
428		virus type 1 and type 2 (HTLV-1 and HTLV-2) in patients infected with human
429		immunodeficiency virus type 1 (HIV-1). Braz J Infect Dis 21:297-305.
430		https://doi.org/10.1016/j.bjid.2017.02.005.
431	32.	Jacob F, Magri MC, Costa EAS, Santos-Fortuna E, Caterino-de-Araujo A. 2009.
432		Comparison of signal-to-cutoff values in first, second, and third generation enzyme
433		immunoassays for the diagnosis of HTLV-1/2 infection in "at-risk" individuals from
434		São Paulo, Brazil. J Virol Methods 159:288-290.
435		https://doi.org/10.1016/j.jviromet.2009.03.024.
436	33.	Jacob F, Santos-Fortuna E de los, Azevedo RS, Caterino-de-Araujo A. 2007.
437		Performances of HTLV serological tests in diagnosing HTLV infection in high-risk
438		population of São Paulo, Brazil. Rev Inst Med Trop Sao Paulo 49:361-364.
439		https://doi.org/10.1590/S0036-46652007000600005.
440	34.	Berini CA, Susana Pascuccio M, Bautista CT, Gendler SA, Eirin ME, Rodriguez C,
441		Pando MA, Biglione MM. 2008. Comparison of four commercial screening assays
442		for the diagnosis of human T-cell lymphotropic virus types 1 and 2. J Virol Methods
443		147:322-327. https://doi.org/10.1016/j.jviromet.2007.09.012.
444	35.	Glas AS, Lijmer JG, Prins MH, Bonsel GJ, Bossuyt PMM. 2003. The diagnostic

19

4	45		odds ratio: a single indicator of test performance. J Clin Epidemiol 56:1129-1135.
4	46		https://doi.org/10.1016/S0895-4356(03)00177-X.
4	47	36.	Sackett DL, Straus S. 1998. On some clinically useful measures of the accuracy of
4	48		diagnostic tests. ACP J Club 129:A17-19. https://doi.org/10.7326/ACPJC-1998-129-
4	49		2-A17.
4	150	37.	Santos FL, Celedon PA, Zanchin NI, Souza WV, Silva ED, Foti L, Krieger MA,
4	151		Gomes Y de M. 2017. Accuracy of chimeric proteins in the serological diagnosis of
4	152		chronic Chagas disease - a Phase II study. PLoS Negl Trop Dis 11:e0005433.
4	153		https://doi.org/10.1371/journal.pntd.0005433.
4	154	38.	Kapprell H-P, Stieler M, Oer M, Goller A, Hausmann M, Schochetman G, Devare
4	155		SG, Qiu X. 2010. Evaluation of a new third-generation ARCHITECT rHTLV-I/II
4	156		assay for blood screening and diagnosis. Diagn Microbiol Infect Dis 67:61-69.
4	157		https://doi.org/10.1016/j.diagmicrobio.2009.12.021.
4	158	39.	Malm K, Kjerstadius T, Andersson S. 2010. Evaluation of a new screening assay for
4	159		HTLV-1 and -2 antibodies for large-scale use. J Med Virol 82:1606-1611.
4	60		https://doi.org/10.1002/jmv.21867.
4	61	40.	Qiu X, Hodges S, Lukaszewska T, Hino S, Arai H, Yamaguchi J, Swanson P,
4	62		Schochetman G, Devare SG. 2008. Evaluation of a new, fully automated
4	463		immunoassay for detection of HTLV-I and HTLV-II antibodies. J Med Virol
4	164		80:484-493. https://doi.org/10.1002/jmv.21083.
4	465	41.	Laperche S, Sauleda S, Piron M, Mühlbacher A, Schennach H, Schottstedt V,
4	466		Queirós L, Uno N, Yanagihara K, Imdahl R, Hey A, Klinkicht M, Melchior W,
4	67		Muench P, Watanabe T. 2017. Evaluation of Sensitivity and Specificity Performance
4	68		of Elecsys HTLV-I/II Assay in a Multicenter Study in Europe and Japan. J Clin

469		Microbiol 55:2180-2187. https://doi.org/10.1128/JCM.00169-17.
470	42.	Malm K, Kragsbjerg E, Andersson S. 2015. Performance of Liaison XL automated
471		immunoassay platform for blood-borne infection screening on hepatitis B, hepatitis
472		C, HIV 1/2, HTLV 1/2 and Treponema pallidum serological markers. Transfus Med
473		25:101-105. https://doi.org/10.1111/tme.
474	43.	Gantner P, Velay A, Guigue N, Barth H, Wendling M-J, Delaugerre C, Fafi-Kremer
475		S. 2017. Performance of the Liaison® XL Murex recHTLV-I/II Immunoassay in the
476		Detection of HTLV-1/2 Antibodies in Serum. Clin Lab 63:997-1001.
477		https://doi.org/10.7754/Clin.Lab.2017.161026.
478	44.	Ly TD, Coignard C. 2016. Performance of the LIAISON [®] XL murex recHTLV-I/II
479		assay. J Clin Virol 82:S88-S89. https://doi.org/10.1016/j.jcv.2016.08.176.
480	45.	Costa EAS, Magri MC, Caterino-de-Araujo A. 2011. The best algorithm to confirm
481		the diagnosis of HTLV-1 and HTLV-2 in at-risk individuals from São Paulo, Brazil.
482		J Virol Methods 173:280-286. https://doi.org/10.1016/j.jviromet.2011.02.018.
483	46.	Sabino EC, Zrein M, Taborda CP, Otani MM, Ribeiro-Dos-Santos G, Sáez-Alquézar
484		A. 1999. Evaluation of the INNO-LIA HTLV I/II assay for confirmation of human
485		T-cell leukemia virus-reactive sera in blood bank donations. J Clin Microbiol
486		37:1324-1328. https://jcm.asm.org/content/37/5/1324.long.
487		
488	Fig 1	Flowchart depicting study design in accordance with the Standards for Reporting
489	of Di	agnostic Accuracy studies (STARD) guidelines.
490		
491	Fig 2	Reactivity index of screening assays obtained in positive (red dots) and negative
492	(blue	e dots) plasma samples under HTLV-1/2 WB analysis. The cut-off value was $IR = 1.0$

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and the area delimited by lines represents the indeterminate zone (RI \pm 10%). Numbers shown for each group are representative of geometric means (± 95% CI); AUC (Area Under Curve); Sen (Sensitivity); Spe (Specificity); Acc (Accuracy); LR (Likelihood Ratio); DOR (Diagnostic Odds Ratio).

Fig 3 Analysis of WB-indeterminate samples using PCR as a gold standard. Acc (accuracy); CI (confidence interval); κ (Cohen's Kappa coefficient); PCR (polymerase chain reaction); Sen (sensitivity); Spe (specificity).

517	TABLE 1.	Indeterminate	HTLV	patterns in	n samples	from Brazil
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WB Pattern	n (%)	Gold ^b	Murex ^c	SYM ^d
gd21 alone	7 (25.9)	7/7	6/7	6/7
gd21+p19	7 (25.9)	7/7	7/7	7/7
gd21+synthetic peptides (46I or 46II)	5 (18.5)	4/5	4/5	4/5
Others ^a	8 (29.7)	8/8	8/8	8/8
HGIP (29)	0	-	-	-
N (30)	0	-	-	-
Total	27 (100)	26/27	25/27	25/27

^aOne band each for gd21 plus p19 plus p28, gd21 plus p19 plus p26 plus p28 plus p32, gd21
plus p19 plus p28 plus p36, gd21 plus p19 plus p26 plus p28 plus p36, plus gd21 plus p19
plus p26 plus p28 plus p32 plus p36, p19 plus p21 plus p26 plus p28 plus p32 plus p36 plus
MTA-1 plus pr53, p19 plus p26 plus p28, synthetic peptide 46II alone

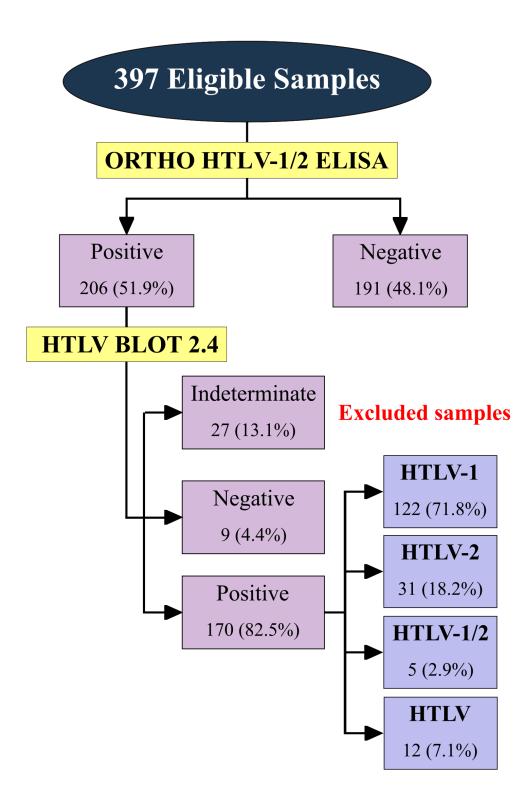
522 ^bGold ELISA HTLV-1/2 (Rem Indústria e Comércio LTDA, São Paulo, Brazil);

⁵²³ ^cMurex HTLV-1/2 (DiaSorin S.p.A., Dartford, UK);

^dAnti-HTLV-1/2 Sym Solution (Symbiosis Diagnostica LTDA, Leme, Brazil)

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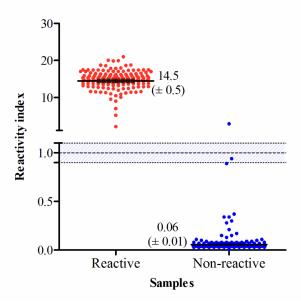
Study design to evaluate the performance of commercial HTLV screening tests

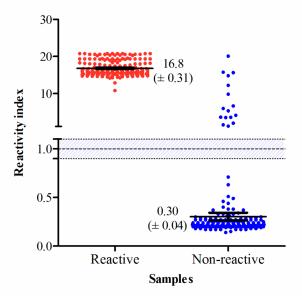


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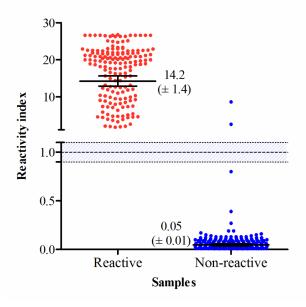
GOLD ELISA HTLV-1/2







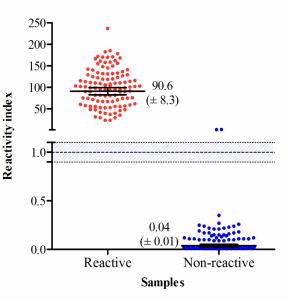
ANTI-HTLV-1/2 SYM SOLUTION



AUC: 0.9989 (± 0.0021) Sen (%): 100 (97.8-100) Spe (%): 99.0 (96.4-99.7) Acc (%): 99.5 (98.1-99.9)

LR+: 100.4 LR-: 0.001 DOR: 168,254

ARCHITECT rHTLV-1/2



AUC: 1.0000	D 1 . 52 4
Sen (%): 100 (96.7-100)	LR+: 52.4
Spe (%): 98.1 (93.3-99.5)	R-: 0.0001
Acc (%): 99.1 (96.7-99.7)	OOR: 524,000



Acc (95%CI)

100

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