

1 Low specificities of HIV diagnostic tests caused by *Trypanosoma brucei gambiense* sleeping
2 sickness

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20 Abstract

21 BACKGROUND: Accuracy of HIV diagnostic tests in tropical infections is poorly
22 documented. Human African trypanosomiasis (HAT) is characterized by a polyclonal B-
23 cell activation, constituting a risk for false positive reactions in diagnostic tests, including
24 HIV tests.

25 METHODS: A retrospective HIV diagnostic test accuracy study was performed on 360
26 human African trypanosomiasis (HAT) patients infected with *T.b. gambiense* before
27 treatment, and 163 patients 2 years after successful treatment in Mbuji Mayi, East Kasai,
28 DR Congo. Sensitivity, specificity and positive predictive value (PPV) of individual tests
29 and algorithms consisting of 3 rapid tests were determined.

30 RESULTS: Sensitivity for all tests was 100% (11/11). Low specificity (96.3%, 335/348) and
31 PPV (45.8%, 11/24) of a classical seroconfirmation strategy (Vironostika ELISA followed
32 by Line Immunoassay) complicated determination of the HIV status, which had to be
33 determined by PCR. Specificities of rapid diagnostic tests were 39.1% for Determine
34 (136/348), 85.3-92.8% (297/348-323/348) for VIKIA, Immunoflow, Doublecheck and
35 Bioline, and 96.6-98.3% (336/348-342/348) for UniGold, Oraquick and STAT-PAK.
36 Specificity for Vironostika was 67.5% (235/348). PPVs ranged between 4.9 and 64.7%.
37 Combining 3 different rapid tests resulted in specificities of 98.3-100% (342-348/348) and
38 PPVs of 64.7-100% (11/17-11/11). In cured HAT patients, specificities were significantly
39 higher for Vironostika, Determine, Unigold and Immunoflow.

40 CONCLUSIONS: *T.b. gambiense* infection decreases the specificity of antibody detection
41 tests for HIV diagnosis. Unless tests have been validated for interference with HAT, HIV
42 diagnosis in untreated HAT using classical algorithms should be avoided. Specific,
43 validated combinations of 3 HIV rapid tests can increase specificity.

44 Introduction

45

46 Over 33 million people live with HIV worldwide, of which two-thirds are in sub-Saharan
47 Africa [16]. Rapid diagnostic tests (RDT) are increasingly used for HIV diagnosis. In 2007,
48 the proportion of RDTs among diagnostic tests procured through World Health Organization
49 (WHO) increased to 96% and the majority were destined to Africa [19]. Accuracy of HIV
50 diagnostic tests in persons suffering from common infections in sub-Saharan Africa is poorly
51 documented. In Tanzania, false positive HIV ELISA results were associated with diagnosis
52 of urinary schistosomiasis and high rheumatoid factor titers [3]. Performance of HIV enzyme
53 immunoassays may be unsatisfactory in visceral leishmaniasis and uncomplicated malaria,
54 resulting in poor positive predictive values [4;12;14]. These false positive results in HIV
55 antibody detection tests have been explained by the polyclonal B-cell activation in visceral
56 leishmaniasis and malaria [4;14].

57 Human African trypanosomiasis (HAT) or sleeping sickness is a fatal disease caused by the
58 protozoan parasites *Trypanosoma brucei (T.b.) gambiense* and *T.b. rhodesiense*, and is
59 transmitted by tsetse flies. In 36 HAT endemic countries in sub-Saharan Africa, about 11000
60 new cases are diagnosed yearly, on a screened population of 2 to 3 million [15]. Polyclonal
61 B-cell activation is also observed in HAT [5]. The uncontrolled antibody production
62 constitutes a risk for false positive reactions in diagnostic tests, including HIV tests, but has
63 hardly been studied, although antibodies cross-reacting with other pathogens have been
64 reported [1;2;13].

65 We assessed the accuracy of commonly used diagnostic tests detecting antibodies against HIV
66 in patients suffering from *gambiense* human African trypanosomiasis and in the cured
67 subgroup of these patients two years after treatment.

68 Materials and methods

69

70 Patients

71 360 human African trypanosomiasis patients infected with *T.b. gambiense* participated to a
72 longitudinal study for monitoring clinical outcomes after HAT treatment, conducted in the
73 hospital of Mbuji Mayi, Kasai Province, DR Congo. Patients and study outcomes are
74 described elsewhere [9]. Patients were prospectively enrolled using the following inclusion
75 criteria 1° trypanosomes in lymph node aspirate, blood or CSF; 2° ≥ 12 years old; and 3°
76 living within a 100 km perimeter around Mbuji Mayi. Exclusion criteria were: 1° pregnancy;
77 2° no guarantee for follow-up; 3° moribund condition; 4° hemorrhagic CSF; 5° concurrent
78 serious illness like tuberculosis, bacterial or cryptococcal meningitis. At time of inclusion, no
79 information was available regarding HIV status. Patients were treated for HAT according to
80 the guidelines of the national program for sleeping sickness of DR Congo. Seventeen patients
81 died during treatment, 14 died during follow-up from non-HAT related reasons, 165 patients
82 experienced treatment failure and 32 did not attend the 24 months visit [9]. Cure of HAT,
83 assessed at 24 months post-treatment, was achieved in 163/360 patients. The Ministry of
84 Health, DR Congo, and the Ethical Committee of the University of Antwerp, Belgium,
85 approved the protocol as well as an amendment allowing for HIV testing. We report on
86 results of HIV tests performed retrospectively at the Institute of Tropical Medicine in
87 Antwerp, on specimens taken before HAT treatment and, for 163 cured patients, also at 24
88 months after HAT treatment.

89

90 Specimens

91 Blood taken on clotting activator was allowed to clot for 1 hour at ambient temperature and
92 centrifuged at 1000g for 15 minutes. Serum was collected and immediately frozen in liquid
93 nitrogen, shipped on dry ice and stored at -80°C until use.

94 Blood taken on heparin was mixed with an equal volume of AS1 storage buffer (Qiagen,
95 Germany). Specimens were shipped and stored at ambient temperature. DNA was extracted
96 with the QIAamp DNA blood mini kit (Qiagen, Germany).

97

98 Reference HIV tests

99 Serum specimens were screened for presence of HIV specific antibodies and antigens using
100 the Vironostika HIV Uni-Form II Ag/Ab ELISA test (bioMérieux, Boxtel, The Netherlands).
101 If the result of Vironostika was reactive (optical density>cut-off) and taking into account a
102 grey zone (cut-off-20%<optical density<cut-off), serum was tested with the INNO-LIA™
103 HIV I/II Score line immuno-assay (Innogenetics, Ghent, Belgium). Following the
104 instructions of the manufacturer, INNO-LIA results were interpreted as negative, positive or
105 indeterminate. INNO-LIA was followed by the INNOTEST® HIV Antigen mAb ELISA
106 (Innogenetics, Ghent, Belgium) to detect early seroconversions. To confirm or exclude HIV
107 infection in certain cases, HIV PCR was performed, following a nested method in an
108 algorithm of three different primer sets in pol, env and LTR region. The obtained sensitivity
109 and specificity of this PCR is 100%. [17].

110

111 Rapid diagnostic tests for detection of HIV specific antibodies

112 All serum specimens were tested in Determine™ HIV 1/2 (Inverness Medical, Unipath Ltd.
113 Bedford, United Kingdom), Uni-Gold™ HIV (Trinity Biotech plc, Bray, Co Wicklow,
114 Ireland), OraQuick Rapid HIV-1/2 Antibody Test (Orasure Technologies Inc., Bethlehem, PA
115 USA), DoubleCheckGold™ HIV 1&2 (Orgenics, Yavne, Israel), SD Bioline HIV-1/2 3.0

116 (Standard Diagnostics Inc., Yongin-si, Kyonggi-do, Korea), ImmunoFlow HIV1-HIV2 (Core
117 Diagnostics, Birmingham, United Kingdom), VIKIA[®] HIV 1/2 (bioMérieux, Marcy-L'Etoile,
118 France) and HIV 1/2 STAT-PAK assay (Chembio Diagnostic systems Inc, Medford, NY
119 USA), following the instructions of the manufacturers.

120

121 Data analysis

122 2x2 tables were constructed. STATA version 10 was used to calculate specificities and
123 positive predictive values (PPV) with 95% confidence intervals (CI). Specificities were
124 compared using the McNemar chi square test.

125 Results

126

127 HIV status of HAT patients before treatment

128 Stored serum was available for 359/360 HAT patients. Determination of the HIV status of the
129 HAT patients was unexpectedly complicated. In Vironostika, 25/359 specimens were
130 classified as grey zone and 99/359 as reactive. In INNO-LIA, 51/124 of these specimens
131 tested indeterminate and 26/124 were positive. Eleven INNO-LIA positive sera reacted
132 strongly with all HIV-1 test lines (sgp120, gp41, p31, p24, and p17). The remaining 15
133 INNO-LIA positive sera, reacted weakly and had unusual, incomplete HIV-1 line
134 combinations (no reaction with p31, p24, or p17 in 8-14 samples). To detect possible
135 seroconverters, all 113 specimens that were Vironostika grey zone or reactive and INNO-LIA
136 negative, indeterminate or weakly positive, were tested in INNOTEST[®] HIV Antigen, and
137 were negative. Facing an unusual high number of unclear results with the classical
138 serological confirmation strategy, PCR was performed on INNO-LIA positives to solve the
139 specificity problem. Only the 11 strong INNO-LIA positives were confirmed to be HIV
140 positive in PCR.

141 Thus, HIV prevalence in HAT patients was 3.1% (11/359, CI 1.5-5.4%). Specificity of
142 Vironostika was 67.5% (235/348, CI 62-72%) with a positive predictive value (PPV) of 8.9%
143 (CI 4.5-15.3%). Considering samples HIV positive if both Vironostika reactive and INNO-
144 LIA positive, specificity and PPV of the classical serological confirmation strategy were
145 respectively 96.3% (335/348, CI 94.2-98.3%) and 45.8% (11/24 CI 25.4-66.3%).

146

147 Results of HIV RDT in HAT patients before treatment

148 Sensitivity of all individual rapid tests was 100% (11/11). Specificities (table 1) were 39.1%
149 for Determine, 96.6% for UniGold, 98.0% for Oraquick, 91.7% for Doublecheck, 92.8% for

150 Bioline, 90.2% for Immunoflow, 85.3% for VIKIA and 98.3% for STAT-PAK. Positive
151 predictive values (table 1) ranged from 4.9 to 64.7%.

152 False positivity seemed randomly scattered with one out of 348 HIV negative samples
153 reactive in 6 RDTs, 2/348 in 5 RDTs, 6/348 in 4 RDTs, 17/348 in 3 RDTs, and 63/348 HIV
154 negative samples reactive in 2 RDTs.

155 HIV diagnosis in low prevalence settings should be based on the combination of 3 different
156 RDT's, considering only those patients HIV positive if 3 assays are reactive. All test
157 algorithms consisting of 3 different RDTs result in 100% sensitivity, specificities ranged
158 between 98.3% (6 false positives, 342/348, 96.9-100%) and 100% (348/348). Seventeen
159 specific three-test combinations provided 100% specificity (Table 2).

160

161 HIV status of cured HAT patients, 2 years after HAT treatment

162 Two years after treatment for HAT, 163/360 patients were declared cured from HAT. From
163 162, serum was available. Among these 162 patients, one was HIV positive before HAT
164 treatment, and was reconfirmed by HIV PCR two years post-treatment. Amongst the 161
165 remaining samples, 4 were grey zone or reactive in Vironostika and indeterminate or positive
166 in INNO-LIA. They were INNOTEST[®] HIV Antigen and PCR negative. Thus all 161 HIV
167 pre-treatment negative patients remained HIV negative. Specificity of the Vironostika test
168 increased significantly (Table 3).

169

170 Results of HIV RDT in cured HAT patients

171 After cure from HAT, the HIV positive patient remained reactive in all HIV RDTs.
172 Specificity increased significantly for Determine, Unigold and Immunoflow tests and was
173 higher but not significant for the Oraquick, Bioline, VIKIA and STAT-PAK tests (table 3).
174 For Doublecheck, specificity was significantly lower on specimens from cured patients.

175 Discussion

176

177 When testing *T.b. gambiense* HAT patients, low specificities were observed in antibody
178 detection RDTs as well as in reference tests (ELISA and Line ImmunoAssay) for HIV
179 diagnosis. On cured HAT patients, specificities were higher. Sleeping sickness thus
180 decreases the specificity of HIV diagnostic antibody detection tests, including tests and
181 algorithms used for confirmation.

182 Test specificities of HIV RDTs observed in HAT patients are below specificities of $\geq 99\%$
183 reported by WHO [18]. Surprisingly, this is the first report on low specificity of HIV tests in
184 *T.b. gambiense* HAT. Other studies examining HIV prevalences in HAT did not report
185 specificity problems of the second generation ELISA and Western Blot tests used [7;8;10;11].
186 False positive results of HIV ELISAs were observed in uncomplicated malaria, with
187 specificities of 96.4 and 98.4% and positive predictive values of 53 and 57% [4] and visceral
188 leishmaniasis with 9% of false positive results [14]. Urinary schistosomiasis and rheumatoid
189 factor were associated with false positive HIV test results [3]. Although *T.b. gambiense* is not
190 the only protozoan parasite impairing HIV test specificity, its effect seems more profound.
191 This might be due to the strong non-specific polyclonal B-cell activation and/or high
192 concentration of rheumatoid factor-like anti-immunoglobulin antibodies occurring in sleeping
193 sickness [5;6]. The specificity differences between tests might be caused by different
194 antigens, or different involvement of IgM. To increase sensitivity, several third generation
195 tests include IgM in the reaction. Such HIV tests are at higher risk for non-specific reactions,
196 as antibodies resulting from the B-cell activation in HAT belong mainly to the IgM class [5].
197 In the absence of an endemic HAT negative control group, the relationship between the low
198 test specificity and HAT could be demonstrated by comparing pre-and post-HAT treatment
199 specificities in the same patients. Presence of other immunological disorders or biological

200 factors such as co-infections in our HAT patients were not examined. Although we cannot
201 exclude their interference in the performance of the HIV tests, this seems unlikely since the
202 same tests were repeated in the same persons after cure from HAT. The overall specificities
203 of HIV RDTs in the region remain unknown, although they are expected to be equal or higher
204 than the observed specificities in cured HAT. The accuracy of DoubleCheckGold™ HIV
205 1&2 should be further investigated, since we have no explanation for its decrease in
206 specificity after successful treatment for HAT, despite use of the same test batch number.

207 Considering the difficulties we encountered at the reference laboratory in determining the real
208 HIV status of HAT patients, it is obvious that the risk for wrong diagnosis is quite high. Not
209 only is there the risk of false positive HIV diagnosis in patients with confirmed HAT, but also
210 of HAT being misdiagnosed as HIV, leaving the HAT patient without treatment. Confounding
211 HAT and HIV diagnosis may become particularly problematic in the context of integration of
212 HAT control in basic health infrastructures [15]. Clinicians easily oversee HAT if they are
213 not familiar with the disease, since the clinical picture may mimic malaria, tuberculosis,
214 toxoplasmosis, viral encephalitis, brucellosis, lymphoma, typhoid fever, neurosyphilis, HIV
215 and opportunistic infections (e.g. cryptococcosis, toxoplasmosis). One should be careful to
216 generalize our observations since we haven't tested the accuracy of HIV antibody tests in *T.b.*
217 *rhodesiense* HAT patients.

218 Our results support the recommendation by WHO to perform 3 different RDTs for HIV
219 diagnosis in settings with prevalences lower than 10% [18]. Considering only those persons
220 HIV positive if they are reactive in 3 different tests, increases the total specificity of HIV
221 RDTs in HAT, although up to 1.7% of false positivity remained. Although we report on
222 specificities of different RDTs and of test algorithms for HIV diagnosis in *T.b. gambiense*
223 HAT patients, we withhold from proposing an optimal test algorithm for *T.b. gambiense*
224 endemic regions. In the present study, the number of HIV positive samples was too low to

225 make reliable estimations of test sensitivities. Performance of the various assays, both
226 reference tests and RDT, remains to be determined in a HAT negative population. Moreover,
227 besides test sensitivity and specificity other factors influence the choice of the most suitable
228 test algorithm, such as cost, test availability, robustness, user friendliness, test rapidity and
229 necessary equipment.

230 In conclusion, *T.b. gambiense* infection decreases the specificity of antibody detection tests
231 for HIV diagnosis. Even a gold standard seroconfirmation strategy does not provide a
232 solution for determining the real HIV status, and limited data on the performance of a
233 Western blot test also show lower specificity caused by indeterminate results. Since the risk
234 for HIV misdiagnosis is considerable, HIV diagnosis should not be established in the setting
235 of an untreated HAT infection using classical algorithms, unless the applied tests have been
236 validated specifically for non-interference with HAT. Although a risk for false positive
237 reactions in HAT remains, specific combinations of 3 different serially applied HIV RDTs
238 increase specificity and should be applied for determination of the HIV status.

239

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308 Table 1: Specificity and positive predictive value (PPV) with 95% confidence intervals (CI)
 309 observed with diagnostic HIV tests in 348 untreated human African trypanosomiasis patients
 310

	Pre-treatment specificity	PPV
	n/N (% , 95% CI)	n/N (% , 95% CI)
Bioline	323/348 (92.8%, 90.1-95.5%)	11/36 (30.6%, 15.2-45.9%)
Determine	136/348 (39.1%, 33.9-44.2%)	11/223 (4.9%, 2.1-7.8%)
DoubleCheck	319/348 (91.7%, 88.7-94.6%)	11/40 (27.5%, 13.4-41.6%)
ImmunoFlow	314/348 (90.2%, 87.1-93.4%)	11/45 (24.4%, 11.7-37.2%)
Oraquick	341/348 (98.0%, 96.5-99.5%)	11/18 (61.1%, 37.9-84.4%)
STAT-PAK	342/348 (98.3%, 96.9-100%)	11/17 (64.7%, 41.2-88.2)
Unigold	336/348 (96.6%, 94.6-98.5%)	11/23 (47.8%, 26.9-68.8%)
VIKIA	297/348 (85.3%, 81.6-89.1%)	11/62 (17.7%, 8.1-27.4%)
Vironostika	235/348 (67.5%, 62.6-72.5%)	11/124 (8.9%, 3.8-13.9%)

311

312

313 Table 2: Combinations of 3 different HIV rapid diagnostic tests providing 100% specificity in
314 348 untreated *T.b. gambiense* patients. Only patients that were reactive in all 3 RDTs were
315 considered HIV positive. The order in which test combinations are presented is alphabetical,
316 reflects parallel testing and is no suggestion for test order in serial testing.

317

Bioline + Determine + STAT-PAK

Bioline + Immunoflow + Oraquick

Bioline + Oraquick + STAT-PAK

Bioline + STAT-PAK + Unigold

Determine + Doublecheck + STAT-PAK

Determine + Immunoflow + STAT-PAK

Determine + STAT-PAK + VIKIA

Doublecheck + Immunoflow + Oraquick

Doublecheck + Immunoflow + Unigold

Doublecheck + Oraquick + STAT-PAK

Doublecheck + STAT-PAK + Unigold

Immunoflow + Oraquick + STAT-PAK

Immunoflow + Oraquick + Unigold

Immunoflow + STAT-PAK + Unigold

Immunoflow + Unigold + VIKIA

Oraquick + STAT-PAK + VIKIA

STAT-PAK + Unigold + VIKIA

318

319 Table 3: HIV ELISA and RDT test results in 161 HIV negative HAT patients, before
 320 treatment for HAT (pre-treatment) and 2 years after successful treatment (post-treatment).
 321 Specificities were compared using the McNemar chi square test.

322

	Pre-treatment specificity	Post-treatment specificity	<i>p</i>
	n/N (% , 95CI)	n/N (% , 95CI)	
Bioline	150/161 (93.2%, 89.2-97.1%)	154/161 (95.7%, 92.5-98.8%)	0.25
Determine	74/161 (46.0%, 38.2-53.7%)	139/161 (86.3%, 81.0-91.7%)	<0.001
DoubleCheck	148/161 (91.9%, 87.7-96.2%)	130/161 (80.7%, 74.6-86.9%)	0.003
ImmunoFlow	147/161 (91.3%, 86.9-95.7%)	155/161 (96.3%, 93.3-99.2%)	0.046
Oraquick	158/161 (98.1%, 96.0-100%)	160/161 (99.4%, 98.2-100%)	0.16
STAT-PAK	158/161 (98.1%, 96.0-100%)	160/161 (99.4%, 98.2-100%)	0.3
Unigold	155/161 (96.3%, 93-99.2%)	160/161 (99.4%, 98.2-100%)	0.03
VIKIA	134/161 (83.2%, 77.4-89.1%)	144/161 (89.4%, 84.6-94.2%)	0.08
Vironostika	119/161 (73.9%, 67.1-80.8%)	158/161 (98.1%, 96.0-100%)	<0.001

323