

RESEARCH ARTICLE

Asymptomatic *Leishmania* infection in HIV-positive outpatients on antiretroviral therapy in Pernambuco, Brazil

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Abstract

Background

Visceral leishmaniasis (VL) in HIV-positive individuals is a global health problem. HIV-*Leishmania* coinfection worsens prognosis and mortality risk, and HIV-*Leishmania* coinfecting individuals are more susceptible to VL relapses. Early initiation of antiretroviral therapy can protect against *Leishmania* infection in individuals living in VL-endemic areas, and regular use of antiretrovirals might prevent VL relapses in these individuals. We conducted a cross-sectional study in Petrolina, Brazil, an VL-endemic area, to estimate the prevalence of asymptomatic *Leishmania* cases among HIV-positive outpatients.

Methods

We invited any HIV-positive patients, aged ≥ 18 -years-old, under antiretroviral therapy, and who were asymptomatic for VL. Patients were tested for *Leishmania* with enzyme-linked immunosorbent assays (ELISA)-rK39, immunochromatographic test (ICT)-rK39, direct agglutination test (DAT), latex agglutination test (KAtex), and conventional polymerase chain reaction (PCR). HIV-*Leishmania* coinfection was diagnosed when at least one VL test was positive.

Results

A total of 483 patients were included. The sample was predominantly composed of single, < 48-years-old, black/*pardo*, heterosexual males, with fewer than 8 years of schooling. The prevalence of asymptomatic HIV-*Leishmania* coinfection was 9.11% (44/483). HIV mono-infected and HIV-*Leishmania* coinfecting groups differed statistically significantly in terms of race ($p = 0.045$), marital status ($p = 0.030$), and HIV viral load ($p = 0.046$). Black/*pardo*

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patients, married patients, and those with an HIV viral load up to 100,000 copies/ml presented higher odds for HIV-*Leishmania* coinfection.

Conclusions

A considerable number of asymptomatic *Leishmania* cases were observed among HIV-positive individuals in a VL-endemic area. Given the potential impact on transmission and health costs, as well as the impact on these coinfecting individuals, studies of asymptomatic *Leishmania* carriers can be useful for guiding public health policies in VL-endemic areas aiming to control and eliminate the disease.

Author summary

Every year up to 90,000 new cases of visceral leishmaniasis (VL) occurs globally. One of the most neglected diseases, VL is endemic in 60 countries in four continents. Brazil is one of seven countries in which 90% of all cases in the world occur. When co-occurring with HIV, VL tends to be more severe than in HIV-negative persons, leading to worse prognosis and frequent relapses. In this study HIV-positive outpatients, aged 18 years or older, under antiretroviral therapy, who were asymptomatic for VL, were tested for *Leishmania*. We observed a prevalence of 9.11% (44/483) of HIV-*Leishmania* coinfection. For those HIV-positive/*Leishmania*-negative individuals who live in VL-endemic areas, early initiation of antiretroviral therapy can be a protective factor against *Leishmania* infection. On the other hand, for HIV-*Leishmania* coinfecting individuals who regularly use antiretrovirals might prevent VL relapses. Studies centering on asymptomatic *Leishmania* carriers can be useful for guiding public health measures in VL-endemic areas seeking to control and eliminate the disease.

Introduction

One of the most neglected diseases [1], visceral leishmaniasis (VL) is a parasitic disease that mostly affects tropical and subtropical regions [2,3]. Endemic to more than 60 countries, it is estimated that 50,000–90,000 cases occur annually worldwide [4]. Brazil and six other countries account for about 90% of all cases [3]. In the American continent, the majority of VL-positive individuals reside in Brazil [5]. Most of VL cases in Brazil are notified in the Northeast region, and Pernambuco is a VL-endemic area [6].

VL is considered to be an opportunistic infection for those living with the human immunodeficiency virus (HIV) [7]. HIV-positive individuals who live in VL-endemic areas have an increased risk of *Leishmania* infection as compared to HIV-negative individuals [8]. When VL occurs in HIV-infected individuals, the prognosis is typically poor and the mortality rate is high [9]. Consequently, several countries have performed studies to estimate the prevalence of *Leishmania* infection in HIV-positive individuals [10–15].

Asymptomatic *Leishmania* infected individuals, despite the typically low parasite load, might contribute to maintaining the transmission cycle of *Leishmania* parasites in endemic regions during episodes of increased parasite load and disease relapse [16,17]. In Brazil, due to the high prevalence of HIV-*Leishmania* coinfection observed in previous studies [10,15,18], and due to the other potential implications (e.g. frequent relapses, mother-to-child

transmission [19]), testing for *Leishmania* should be strongly recommended for all HIV-positive individuals. In addition, once HIV-*Leishmania* coinfection is diagnosed, early start of highly active antiretroviral therapy (HAART) should be recommended as a protective factor against VL relapses [20]. Moreover, for those HIV-positive individuals who are negative for leishmaniasis and are living in VL-endemic areas, HAART could decrease the risk of *Leishmania* infection [21].

The Brazilian national HIV/AIDS program assists all persons who live with HIV with HAART, free of charge, and in the same way it provides treatment for all individuals affected by VL. Despite the impact on the public health system, there have been few studies following up these HIV-*Leishmania* coinfection cases. Most of these studies have focused on hospitalized patients. However, testing for *Leishmania* in HIV-positive outpatients in VL-endemic areas may be useful for guiding health policies aiming to control and eliminate leishmaniasis, and it could improve treatment and outcomes for those affected by the disease.

In this study, we aimed to estimate the prevalence of asymptomatic *Leishmania* cases in HIV-positive outpatients under continuous use of HAART, in a VL-endemic area, Pernambuco, in Northeast Brazil.

Methods

Ethics statement

The study was approved by the research ethics committee of Instituto Aggeu Magalhães, Fiocruz Pernambuco (approval number 51235815.0.0000.5190). All subjects were adults and provided written, informed consent. This study was conducted in accordance with the Declaration of Helsinki.

Study design and sample

We performed a cross-sectional study aiming to estimate the prevalence of HIV-*Leishmania* coinfection in HIV-positive outpatients from the municipality of Petrolina, in the state of Pernambuco, Brazil. Petrolina is an VL-endemic area from where most VL cases in the state of Pernambuco are reported [22]. The study was conducted in a public HIV outpatient clinic, which serviced about 600 individuals regularly using HAART at the time of this investigation.

The study population included any HIV-positive patients, using HAART, aged 18-years-old or older. Individuals under treatment for VL and those with current VL symptoms were excluded. Based on a previous prevalence study in Pernambuco [18], the minimum sample size was calculated to be 159 individuals for a 95% confidence interval (Epi Info 7.2.3.0 software, <https://www.cdc.gov/epiinfo/index.html>). We publicly invited patients at this outpatient clinic to participate in the study. Many of them demonstrated interest in knowing their serological status for *Leishmania*, as they resided in an endemic area. Due to the high demand, we decided to include anyone who requested to be tested. Therefore, there was a higher number of participants recruited compared to the originally calculated sample size. Each participant received the results of the tests done in the study.

Data collection and laboratory procedures

After the interview and physical examination, peripheral venous blood and urine samples were collected from the patients at the same time when the samples were taken for analyzing the lymphocyte T CD4+ (LTCD4+) count or HIV viral load. The samples were stored, processed, and analyzed at Fiocruz Pernambuco, a referral public research center. Participants were tested for VL with enzyme-linked immunosorbent assays (ELISA)-rK39,

immunochromatographic test (ICT)-rK39, direct agglutination test (DAT), latex agglutination test (KAtex) and polymerase chain reaction (PCR) test. All HIV-positive individuals with at least one positive test for VL were considered HIV-*Leishmania* coinfection cases. Hemogram, biochemistry, LTCD4+ count, and HIV viral load data were obtained from the medical records.

For the ELISA-rK39 assays, the commercial recombinant rK39 antigen was purchased from Rekom Biotech (Granada, Spain) and the assays were essentially carried out as previously described by Scalone et al [23] and Abass et al [24]. For ICT-rK39, we used the OnSite Leishmania IgM/IgG Combo test (CTK Biotech, Inc., Poway, CA, USA) following the manufacturer's instructions.

For DAT, we used a freeze-dried antigen from Biomedical Research (Amsterdam, The Netherlands) and titers of 1:3,200 or higher were considered to indicate a positive test [25]. *Leishmania* antigen was detected in urine by means of the KAtex kit (Kalon Biological Ltd., Guildford, UK) according to the manufacturer's instructions.

For PCR, we targeted the kinetoplast DNA of *Leishmania* (kDNA). We used the following primers: 150 5'-GGG(G/T)AGGGGCGTTCT(C/G)CGAA3' and 152 5'-(C/G)(C/G)(C/G)(A/T)CTAT(A/T) TTACACCAACCCC-3', which amplify a fragment of 120 bp for all *Leishmania* species. Details on the PCR conditions were described by Souza et al [26]. To confirm the species (*L. Infantum*), we used primers RLC2 5'-GGGAAATTGGCCTCCCTGAG-3' and FLC2 5'-GTCAGTGTCGGAACTAATCCGC-3', which amplify a product of 230 bp, according to Gualda et al [27]. The results were analyzed by electrophoresis in 1.5% agarose gels stained with ethidium bromide and were visualized under ultra-violet light.

Statistical analysis

Data were entered and stored on spreadsheets using Microsoft Excel Professional Plus 2016 software (Microsoft Corp., Redmond, WA, USA). Data analysis was performed in Stata SE 12.0 software for Windows (StataCorp, College Station, TX, USA).

Frequencies and means with 95% confidence interval of the variables of interest were obtained. We compared an HIV mono-infected and an HIV-*Leishmania* coinfecting group. For binary/categorical variables, the chi-square test was used (significance level $p < 0.05$). For continuous variables, we used the two-sample Wilcoxon rank-sum (Mann-Whitney) test. Variables that yielded a p -value ≤ 0.1 in univariate analysis were included in a multivariate analysis using logistic regression.

Results

Of 487 HIV-positive individuals who agreed to enroll in the study, three individuals were excluded as they presented typical symptoms of VL and one individual was excluded for not using HAART. The study population was composed mainly of single (45.5%), black/*pardo* (83.4%), heterosexual (58.2%) men (61.3%) with 8 or fewer years of schooling (48.24%). About 3% reported using intravenous drugs (Table 1).

The prevalence of HIV-*Leishmania* coinfection was 9.11% (44/483). The highest positivity by VL test was seen with DAT (3.53%), followed by ELISA-rK39 (2.48%), and PCR kDNA (2.28%) (Table 2). Two individuals tested positive with DAT and KAtex, and one person tested positive with DAT and rK39. In addition, of the individuals who tested positive for *Leishmania* spp, three (6.8%) reported having had previous diagnoses of VL, and all of whom reported having been treated.

Regarding general laboratory findings, all results were compatible with the reference standards (Table 3). When the two groups were compared, we observed statistically significant

Table 1. Comparison of epidemiological characteristics between HIV-*Leishmania* coinfecting and HIV mono-infected groups (univariate analysis) among HIV-positive individuals tested for *Leishmania* in Petrolina, Brazil.

Variables	HIV mono (N = 439)		HIV- <i>Leishmania</i> (N = 44)		p-value
	n	(%)	n	(%)	
Gender					
Female	170	(38.7)	17	(38.6)	
Male	269	(61.3)	27	(61.4)	0.991
Age (years)					
18–27	70	(15.9)	7	(15.9)	
28–37	109	(24.8)	7	(15.9)	
38–47	140	(31.9)	17	(38.6)	0.582
48 or more	120	(27.3)	13	(29.5)	
Years of schooling					
0–8	211	(48.1)	22	(50.0)	0.751
9–11	162	(36.9)	14	(31.8)	
12 or more	66	(15.0)	8	(18.2)	
Marital status					
Separated/divorced/widowed	65	(14.8)	2	(4.5)	
Married/stable union	168	(38.3)	28	(63.6)	0.003
Single	206	(46.9)	14	(31.8)	
Race					
White	73	(16.6)	1	(2.3)	
Black/ <i>pardo</i>	336	(83.4)	43	(97.7)	0.039
Indigenous	1	(0.2)	0	(–)	
Sexual orientation					
Did not inform	86	(19.6)	7	(15.9)	
Heterosexual	257	(58.5)	24	(54.5)	0.576
Homosexual	76	(17.3)	9	(20.4)	
Bisexual	20	(4.6)	4	(9.1)	
Use of intravenous drugs					
No	426	(97.0)	43	(97.7)	0.795
Yes	13	(3.0)	1	(2.3)	
Dogs at home					
No	274	(91.3)	165	(90.2)	0.665
Yes	26	(8.7)	18	(9.8)	

All percentages are column percentages

Pardo is a specific Brazilian self-declared race, non-white

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Table 2. Prevalence of asymptomatic HIV-*Leishmania* coinfection cases in outpatients from Petrolina, Brazil, according to VL test done.

VL tests	Positivity	Prevalence (%)
ELISA-rK39	12/483	2.48
rK39-ICT	5/470	1.06
DAT	17/482	3.53
KAtex	2/483	0.41
PCR kDNA	11/482	2.28
DAT and KAtex	2/482	0.41
DAT and rK39-ICT	1/470	0.21
Total (at least one positive test)	44/483	9.11

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Table 3. Comparison of general laboratory characteristics between HIV-*Leishmania* coinfecting and HIV mono-infected groups (univariate analysis) among HIV-positive individuals tested for *Leishmania* in Petrolina, Brazil.

Variables	HIV mono		HIV- <i>Leishmania</i>		p-value
	Mean	(95CI)	Mean	(95CI)	
White blood cells (cells/mm ³)	5793	(5583–6002)	5325	(4770–5881)	0.289
Hemoglobin (g/dL)	13.8	(13.6–14.0)	13.7	(13.0–14.4)	0.706
Platelets (x10 ³) (cells/mm ³)	255	(247–264)	263	(238–288)	0.586
AST (U/L)	31.3	(29.1–33.5)	31.1	(26.4–36.0)	0.880
ALT (U/L)	29.6	(27.7–31.5)	26.8	(22.1–31.4)	0.918
Creatinine (mg/dL)	0.82	(0.80–0.85)	0.90	(0.82–0.97)	0.188

95CI, 95% confidence interval; AST, aspartate aminotransferase; ALT, alanine aminotransferase

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differences in terms of marital status ($p = 0.003$) and race ($p = 0.039$), with HIV-*Leishmania* coinfection being more frequent in married and black/*pardo* individuals. No general laboratory characteristic showed a statistically significant difference between the groups. In terms of HIV infection status (Table 4), the LTCD4+ count was greater than 350 cells/mm³ in 74.4% of the general sample, with no difference between the two groups. Most individuals (73.4%) had an undetectable HIV viral load. When we correlated the LTCD4+ count with the VL tests performed, we observed no statistically significant differences.

Marital status, race, and HIV viral load remained significant in the multivariate model (Table 5). Black/*pardo* individuals were at an increased risk (odds ratio, OR: 7.85; $p = 0.044$) of being HIV-*Leishmania* co-infected, as compared with white individuals. Marriage/stable unions and a detectable HIV viral load up to 100,000 copies/mL were also associated with an increased risk (OR: 5.12, $p = 0.029$ and OR: 2.01, $p = 0.047$, respectively).

Discussion

This study focused on HIV-*Leishmania* coinfection in outpatients in Pernambuco, which had not been reported previously. The prevalence of asymptomatic HIV-*Leishmania* coinfection was 9.11% (44/483). There were statistically significant differences between the HIV mono-infected and HIV-*Leishmania* coinfecting groups in term of race ($p = 0.045$), marital status ($p = 0.030$), and HIV viral load ($p = 0.046$). Black/*pardo* patients, married patients, and those

Table 4. Comparison of HIV infection-related characteristics between HIV-*Leishmania* coinfecting and HIV mono-infected groups (univariate analysis) among HIV-positive individuals tested for *Leishmania* in Petrolina, Brazil.

Variables	HIV mono		HIV- <i>Leishmania</i>		p-value
	n	(%)	n	(%)	
HIV viral load (copies/ml)					
Undetectable (<50)	326	(74.4)	28	(63.6)	0.099
Up to 100,000	90	(20.5)	15	(34.1)	
100,000 or more	22	(5.0)	1	(2.3)	
LTCD4+ count (cells/mm³)					
Up to 200	43	(10.1)	5	(11.9)	
200 to 350	66	(15.5)	6	(14.3)	
350 or more	318	(74.5)	31	(73.8)	0.922

All percentages are column percentages; LTCD4+, lymphocyte T CD4+

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Table 5. Odds ratios for asymptomatic HIV-*Leishmania* coinfection based on multivariate analysis.

Variable	OR	95CI	p-value
Race			
White	1		
Black/ <i>pardo</i>	7.85	1.05–58.39	0.044
Marital status			
Separated/Divorced/Widow	1		
Married/Stable union	5.12	1.17–22.29	0.029
Single	1.99	0.44–9.09	0.374
HIV viral load (copies/ml)			
Undetectable (<50)	1		
Up to 100,000	2.01	1.01–4.05	0.047
100,000 or more	0.70	0.09–5.60	0.740

OR odds ratio; 95CI 95% confidence interval

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with an HIV viral load up to 100,000 copies/mL presented higher odds for HIV-*Leishmania* coinfection.

The prevalence of asymptomatic HIV-*Leishmania* coinfection in this study (9.1%) was lower than that in a previous study in Pernambuco of hospitalized HIV-positive patients who were tested for VL (16.9%) [18]. This previous study involved three referral hospitals for infectious diseases that serviced the entire state. Despite the higher percentage, compared with the current study, there were fewer cases (35 vs 44) and we tested more individuals (483 vs 207). A similar study in Minas Gerais, Southeastern Brazil, observed a prevalence of asymptomatic *Leishmania* infection in HIV-positive individuals of 20% [15]. Minas Gerais is a Brazilian state with high VL endemicity, and it would be expected to have a higher prevalence than that observed in our study. In the Metema district of Northwestern Ethiopia, a pilot study in HIV-infected adults identified a prevalence of 12.8% in males and 4.2% in females for asymptomatic HIV-*Leishmania* coinfection cases, based on the same tests used in the present study [28].

In the VL tests used, we observed a low coincidence of results. Based on previous studies [15,28,29], a large variation in the test results could be expected. Furthermore, the sample consisted of HIV-positive individuals with no suspicion of VL. The highest prevalence rate was observed with the DAT, a test with high sensitivity that is considered to be a good diagnostic tool for immunocompromised individuals [30,31]. The DAT has also showed the highest prevalence rate in a previous prevalence study in Pernambuco with symptomatic patients [18]. We observed a low prevalence with the ICT-rK39 rapid test (1%). Apparently, this is not the best serological screening test for individuals living in *L. infantum*-endemic regions [15,18,29,30]. Despite the reduced sensitivity of serological tests for VL in HIV-positive individuals [8,30], this type of test should not be excluded, as a positive result should be considered when associated with clinical features [32]. Serological tests are important as screening tests for VL, particularly in VL-endemic areas and for detection of asymptomatic HIV-*Leishmania* coinfecting persons, as they normally have higher LTCD4+ counts. In our study, most participants had an LTCD4+ count exceeding 350 cells/mm³, and thus they had potentially similar humoral response as HIV-negative persons.

In terms of epidemiological aspects, we observed statistically significant differences for race and marital status. Indeed, in Brazil race and poverty are strongly connected, and most people affected by neglected tropical diseases live in low income regions [33], such as the region in which this study was performed.

The mean LTCD4+ count in HIV-*Leishmania* co-infected individuals was similar to that observed in the HIV mono-infected group. In the present study, almost three-quarters of all individuals presented with LTCD4+ exceeding 350 cells/mm³. Most participants from the previous study of hospitalized patients in Pernambuco had LTCD4+ counts lower than 200 cells/mm³ in both groups (VL-HIV and HIV) [18]. New cohort studies with paired samples may better explain the behavior of *Leishmania* infection in our population. In the present study, all HIV-positive individuals were on HAART, which explains the higher LTCD4+ count, and which might be a protective factor against developing VL.

We observed that, a detectable viral load, although lower than 100,000 copies/ml, was associated with *Leishmania* infection. As all patients were asymptomatic, this could indicate an initial HAART failure or irregular use of the treatment. In the previous study of hospitalized VL-HIV coinfecting patients, only 16% had an undetectable viral load [18], while in the present study the viral load was undetectable in 63% of the coinfecting individuals. Since regular use of HAART usually increases the LTCD4+ count, which, in turn, is a protective factor against VL relapses, campaigns to encourage the regular use of antiretrovirals should be intensified among individuals living with HIV in VL-endemic areas. In Brazil, to date, only secondary prophylaxis for VL is recommended, and the only one marker used to guide this prophylaxis is the LTCD4+ count.

The factors determining maintenance of an asymptomatic VL state have not yet been established. This balance between the parasite infection and the host's immune response, such as, for example, in blood donors or in HIV-AIDS patients, probably extends beyond nutritional status and genetic factors. Due to the increased risk of relapses and the poor prognosis, it is important for HIV-positive persons, particularly those living in VL-endemic areas, to know about a previous *Leishmania* infections. Identification of new markers or tests that might delimit active disease, suggest cure, and predict relapses is urgent. It may be challenging to distinguish active VL cases from another opportunistic infection, and new and less-invasive markers for VL could help health professionals in making more accurate diagnoses and consequently avoiding unnecessary treatments.

Conclusions

Visceral leishmaniasis remains an important problem in Brazil, particularly in the Northeastern region. We observed a considerable number of asymptomatic *Leishmania* cases in HIV-positive individuals. Studies focusing on health care of asymptomatic individuals could be useful for public health policies in VL-endemic areas, facilitating monitoring of the progress of leishmaniasis control. In addition, at the individual level, it is important to follow up all these HIV-*Leishmania* coinfecting persons in terms of VL prophylaxis and treatment, and to predict relapses. Consequently, we strongly recommend testing for *Leishmania* in all HIV-positive individuals in VL-endemic areas.

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