Research Article

Studies on the Feeding Habits of Lutzomyia (Lutzomyia) longipalpis (Lutz & Neiva, 1912) (Diptera: Psychodidae: Phlebotominae) Populations from Endemic Areas of American Visceral Leishmaniasis in Northeastern Brazil

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The aim of this study was to identify potential blood feeding sources of L. (L.) longipalpis specimens from populations in Northeastern Brazil, endemic areas of American Visceral Leishmaniasis (AVL) and its correlation with the transmission of L. (L.) i. chagasi. The ELISA technique was applied using bird, dog, goat, opossum, equine, feline, human, sheep, and rodent antisera to analyze 609 females, resulting in an overall positivity of 60%. In all municipalities, females showed higher positivity for bird followed by dog antiserum and sand fly specimens were also positive for equine, feline, human, sheep, goat, opossum, and rodent antiserum. The finding for 17 combinations of two or three types of blood in some females corroborates the opportunistic habit of this sand fly species. The results demonstrating the association between L. (L.) longipalpis and opossum suggest the need for further evaluation of the real role of this synanthropic mammal in the eco-epidemiology of AVL.

1. Introduction

American visceral leishmaniasis (AVL) is a serious public health problem in Brazil and presents a new epidemiological profile associated with domestic environments and, in this context, Lutzomyia (Lutzomyia) longipalpis is important considering its capacity to adapt to a wide range of impacted habitats, in addition to its sylvatic origin [1–3]. The enzyme immunoassay (ELISA) has been used to identify the feeding habits of L. (L.) longipalpis [4–6].

In light of this, studies related to feeding habits of sand fly vector L. (L.) longipalpis could contribute to a better understanding of eco-epidemiology of AVL, discussing its close association with Leishmania (Leishmania) infantum chagasi reservoirs.

Currently, Northeastern Brazil accounts for about 47% of human cases for AVL exhibiting both epidemiological profiles, rural and urban, with highest incidences of the disease according to the Brazilian National Leishmaniasis Program (NLP) [1]. The aim of this study was to identify potential blood meal sources for L. (L.) longipalpis from some Northeastern Brazil endemic municipalities.
2. Materials and Methods

2.1. Sand Fly Precedence. Sand flies were collected from the municipalities of Jequié (State of Bahia, BA), Sobral and Massapê (State of Ceará, CE), and Teresina (State of Piauí, PI). These municipalities were selected based on their levels and profiles of AVL transmission according to NLP: Teresina, urban and intense transmission; Sobral, rural and intense transmission; Jequié, rural and moderate transmission, and Massapê, rural and sporadic transmission [1]. Sand flies were collected from March 2006 to September 2007 during months of high frequency of sand flies.

2.2. Sand Fly Sampling. Sand flies were collected using unbaited modified CDC light traps [7], HP model [8], between 6:00 PM and 6:00 AM For each location (municipality), the same number of site collections was selected (four houses with human cases of AVL and with environmental characteristics adequate for breeding sand flies). Collections were done in the peridomicile; two traps were randomly placed at a minimum distance of 50 meters from the residence and from each other. The collected sand flies were screened in a cold chamber (cryolyzer), and females were stored in plastic microtubes and placed in a −20°C freezer for future analysis.

2.3. Bloodmeal Analysis. L. (L.) longipalpis engorged females or those with some residual blood meal were tested by ELISA, following the method of Burkom et al. [9] modified by Duarte [10] using bird, dog, goat, opossum, equine, feline, human, sheep, and rodent antisera; male specimens of L. (L.) longipalpis were used as a control; they were treated in the same manner as the overfed female samples. The males for the negative control were chosen because they do not feed on blood and were used to pinpoint false positives in the samples of female sand flies. The choice of antisera was based on the most common animals at collection sites. The L. (L.) longipalpis specimens were identified according to diagnostic morphological features (cibarium and spermatheca) [11].

To the analysis, the abdomen of the frozen female specimens, as well as the controls (males), were macerated in PBS (pH 7.2, 0.001 M) and kept at −20°C until processing. Samples were diluted 1:20 in carbonate bicarbonate buffer (pH 9.6; 0.05 M, Sigma) and applied to 96-well polystyrene microplate (NuncC, 442404, maxisorp, Denmark). After incubation (at 37°C for 2 h), plates were washed in PBS/Tween 20–0.05% (Sigma Chemicals Co-St. Louis, USA). The next steps involved adding antiserum (PBS/Tween 20 plus 1% skim milk—Molico-Nestlé, Brazil) into the wells and incubating the microplate at 37°C for 30 min (goat anti-rabbit serum peroxidase conjugate—Sigma Chemical USA). Should be washed, the diluted conjugate, at 1:20,000 was added, and after another incubation and wash, the developing buffer (citrate/phosphate pH 5.0–0.05 M hydrogen peroxide (H₂O₂) 30 vol. Merck Diagnostica-RJ, Brasil and O-Phenylenediamine—OPD, Sigma Immunochemicals Co. USA) was applied. The reaction was stopped after 15 min by adding 50 µL of sulfuric acid solution 1 N and read on an ELISA plate reader (WI, USA) using 490 nm operational and 630 nm reference filters.

In every plate, positive controls were used consisting of homologous sera diluted in carbonate bicarbonate buffer (pH 9.6, 0.05 M, Sigma) 200 times. For validation, amounts over 1.0 were expected. In order to estimate the positivity of the samples, a calculation was made based on the average of the absorbance obtained from the reactions observed in heterologous sera plus two standard deviations (cutoff point). This procedure was adopted to exclude the results of possible crossover reactions and to increase the specificity of the assay.

The antisera used were obtained from the Immunodiagnostics Laboratory, Department of Biological Science, Escola Nacional de Saúde Pública Sérgio Arouca, FIOCRUZ; because there are no antibodies to all food sources, we used anti-total protein, which can increase the sensitivity of the method, since the insect feeds on blood. The assay sensibility is estimated (96%) in the Standard Operational Procedure described in the Laboratory where the analyses were performed and described in the reference Duarte [10].

3. Results

A total of 609 sand fly females were analyzed and a general reactivity index of 60% was obtained; specimens from Massapê displayed the greatest diversity in feeding sources, 19 antisera or combinations of antisera unlike the population from Jequié, which was reactive to only six. The percentages of positivity to the antisera tested in four populations of L. (L.) longipalpis are presented in Table 1.

Bird, dog, and equine antisera were positive for all L. (L.) longipalpis populations, with highest positivity observed in blood of birds (Jequié, 36.0%, Teresina, 67.6%, Sobral, 29.9%, Massapê, 51.0%) followed by blood of dogs (Jequié, 16.0%, Teresina 9.5%, Sobral 14.3%, Massapê 2.3%). Equine antisera were reactive in all populations (Jequié, 16.0%, Teresina 2.7%, Sobral 15.6%, Massapê 8.5%). Blood from opossum was detected in L. (L.) longipalpis females from three populations (Jequié, 8.0%, Sobral, 2.6%, and Massapê, 2.3%), and blood of sheep has also been identified in three populations (Teresina 2.7%, Sobral 9.1%, and Massapê 10.8%). Human antiserum was reactive in L. (L.) longipalpis females from Jequié (20%) and Massapê (0.8%). Feline and goat antisera showed positivity for females from Sobral and Massapê, respectively. And rodent antisera were reactive only in Massapê species.

Females that had fed on more than one blood meal source for all L. (L.) longipalpis populations were detected; especially those from Sobral and Massapê, specimens were found that had fed on at least three distinct sources. From the 17 blood combinations detected, dog + bird, dog + sheep, dog + equine, dog + opossum, sheep + feline, and goat + sheep + equine were the most frequent.

4. Discussion

Previous reports based on field studies have suggested that L. (L.) longipalpis has a varied diet, feeding on a wide
Table 1: Percentage of positive females of *Lutzomyia (Lutzomyia) longipalpis.* analyzed by ELISA collected in the Municipality of Jequié (BA), Teresina (PI), Sobral (CE) and Massapê (CE), North-east Brazil.

| Antisera reactivity % | Jequié (BA) | Teresina (PI) | Sobral (CE) | Massapê (CE) |
|-----------------------|-------------|---------------|-------------|--------------|
| Bird                  | 36.0        | 67.6          | 29.9        | 51.0         |
| Dog                   | 16.0        | 9.5           | 14.3        | 2.3          |
| Goat                  | —           | —             | —           | —            |
| Opossum               | 8.0         | —             | 2.6         | 2.3          |
| Equine                | 16.0        | 2.7           | 15.6        | 8.5          |
| Feline                | —           | —             | 1.3         | —            |
| Human                 | 20.0        | —             | —           | 0.8          |
| Sheep                 | —           | 2.7           | 9.1         | 10.8         |
| Rodent                | —           | —             | —           | 0.8          |
| Dog + Human           | 4.0         | —             | —           | —            |
| Dog + Bird            | —           | 4.1           | —           | 2.3          |
| Dog + Sheep           | —           | 8.1           | 3.9         | 4.6          |
| Dog + Equine          | —           | —             | 7.8         | 3.9          |
| Equine + Sheep        | —           | —             | 2.6         | —            |
| Goat + Equine         | —           | —             | 2.6         | —            |
| Dog + Opossum         | —           | —             | 1.3         | 3.1          |
| Sheep + Feline        | —           | —             | 1.3         | 0.8          |
| Bird + Equine         | —           | —             | 1.3         | —            |
| Bird + Opossum        | —           | —             | —           | 1.5          |
| Bird + Sheep          | —           | —             | —           | 0.8          |
| Human + Sheep         | —           | —             | —           | 0.8          |
| Dog + Human + Equine  | —           | —             | —           | 1.5          |
| Bird + Dog + Equine   | —           | —             | —           | 0.8          |
| Bird + Opossum + Dog  | —           | —             | —           | 0.8          |
| Goat + Sheep + Equine | —           | —             | 5.2         | 2.3          |
| Dog + Sheep + Bird    | —           | 5.4           | —           | —            |

range of animals, including dogs, pigs, horses, cattle, and chickens [3, 12–14]. Along with favorable environmental conditions, it has been suggested that population growth of this sand fly vector is determined by the abundance of food sources, which facilitates its adaptation to human dwellings, especially in rural areas.

The data from the present study show that specimens from the four analyzed populations fed mainly on birds, based on the significant percentage of positivity compared to other feeding sources. The strong attraction to birds has already been observed in many other studies in Brazilian States such as Maranhão (MA) [15–17] and Mato Grosso do Sul (MS) [5]. It is known that when chicken coops are in a peridomicile, they attract sand flies and can act as breeding sites. This proximity increases contact between vectors and humans, suggesting that chicken coops play an important epidemiological role [18]. Although chickens may be refractory to *Leishmania* infection; they serve as a feeding source for sand flies and attract potential reservoirs of *L. (L.) i. chagasi*, allowing the establishment and maintenance of AVL transmission in rural areas [19]. In Argentina, a study on the spatial distribution of *L. (L.) longipalpis* found a positive association between the vector and the presence of chicken coops and could contribute to the design of control strategies, defining priority areas for prevention and control [20].

The positivity to dog blood was confirmed for the *L. (L.) longipalpis* populations analyzed. Since the pioneering studies on epidemiology of AVL in Brazil (Sobral/CE), the attraction to dogs was evident [12]. In most AVL transmission areas, there is epidemiological evidence implicating dogs as domestic reservoirs [3]. Studies have been suggesting that dogs are an important blood source for *L. (L.) longipalpis* in Teresina (PI) and Araçatuba (SP), both areas for high incidence of AVL [21, 22]. However, studies conducted in Campo Grande (MS), an area with high prevalence of AVL human cases and where the presence of dogs was observed in all domestic sites, showed a low reactivity for dog blood [5]. In this way, sand fly populations may present different feeding preferences in different ecotypes.

A low positive reaction to feline antiserum was only found in *L. (L.) longipalpis* females from Sobral and Massapê. In fact, there is no evidence for a role of felines in the AVL epidemiology, although recently the experimental infection of *L. (L.) longipalpis* by *L. infantum (= L. infantum chagasi)* was described after xenodiagnosis on a naturally infected cat from an endemic area in Belo Horizonte (MG) [23].

In the present study, even with low indices of positivity, opossum blood in specimens from three populations was detected: Sobral, Massapê, and Jequié. This is the first report of an association between *L. (L.) longipalpis* and opossums in Brazil. In Colombia, a low reactivity for opossums was also observed in *L. (L.) longipalpis* [24]. These data should be analyzed carefully since the opossum (*Didelphis albiventris*) has been suggested as a possible secondary reservoir host for *L. (L.) i. chagasi*; based on studies conducted in Jucá/BAR, only 2 of 84 animals were positive, a result that for some authors has no significant epidemiological importance [25, 26]. Subsequently, other studies have reported natural infections of *Didelphis marsupialis* by *Leishmania* spp., possibly *L. (L.) i. chagasi*, arguing the role of these mammals as potential AVL reservoirs [27, 28]. In light of the evidence for natural infection in opossums and the finding that these animals are feeding sources for *L. (L.) longipalpis*, the possibility that this synanthropic mammal participates in the transmission cycle of AVL in some regions should be considered, especially those that have been undergoing environmental alterations, which facilitates contact of these animals with human habitation.

The anthropophily of *L. (L.) longipalpis* has been observed in fieldworks [1–3, 12, 19], which is one essential criteria in implicating sand fly species as vector. Otherwise, studies evaluating the blood meal of this sand fly conducted in Marajó Island (State of Pará) and in Campo Grande (MS)
demonstrated its high anthropophily [4, 5]. Only two L. (L.) longipalpis populations (Jequié and Massapê) in this study contained human blood in low percentage. Similar results obtained in Araçatuba suggested low anthropophily of L. (L.) longipalpis [22], which was also observed in studies conducted in Colombia [24]. The abundance of feeding sources (birds and dogs) surrounding houses should be considered, as observed at the collection sites in Teresina and Sobral, it can be a determining factor in the choice of the insect’s feeding source, corroborating the opportunistic behavior of this sand fly.

The finding of more than one blood source in some females, mainly in Massapê (up to three), is a strong evidence of the eclectic diet of L. (L.) longipalpis. This behavior is common in sand flies, which is to “try out” different hosts until completing their blood meal. This is, undoubtedly, an important aspect in leishmaniases transmission.

The adaptation of L. (L.) longipalpis to different habitats, even its urbanization, remains a great challenge for AVL control. Clearly, the results of this study indicate the eclecticism of L. (L.) longipalpis with regard to blood feeding. In light of entomological surveillance of leishmaniases, information about the biology of the vector, and especially its interaction with reservoir hosts in association with environmental factors, can indicate changes in the transmission profile of AVL and the process of geographical expansion [29]. In this context, the ability to feed frequently on domestic animals such as birds (chickens), dogs and even synanthropic mammals (opossums) is an important attribute that allows L. (L.) longipalpis to maintain AVL transmission in rural environments and the expansion of the disease to urban areas, contributing to the maintenance of two transmission profiles currently found in Brazil.

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References

[1] Brasil, Ministério da Saúde, Secretaria de Vigilância em Saúde, “Manual de Vigilância e Controle da Leishmaniose Visceral Americana,” 2006, Departamento de Vigilância Epidemiológica, Ministério da Saúde, Secretaria de Vigilância em Saúde, Departamento de Vigilância Epidemiológica, 1st edn, Brasília: Editora do Ministério da Saúde, p.120.
[2] R. Lainson and J. J. Shaw, “New Leishmaniases—the Neotropical Leishmania species,” in Microbiology Infections, chapter 13, pp. 241–266, Topley & Wilson’s, 9th edition, 1998.
[3] R. Lainson and B. F. Rangel, “Lutzomyia longipalpis and the eco-epidemiology of American visceral leishmaniasis, with particular reference to Brazil—a review,” Memorias do Instituto Oswaldo Cruz, vol. 100, no. 8, pp. 811–827, 2005.
[4] R. J. Quinnell, C. Dye, and J. J. Shaw, “Host preferences of the phlebotomine sandfly Lutzomyia longipalpis in Amazonian Brazil,” Medical & Veterinary Entomology, vol. 6, no. 3, pp. 195–200, 1992.
[5] A. G. de Oliveira, A. M. Marassá, C. A. Consales et al., “Observations on the feeding habits of Lutzomyia longipalpis (Lutz & Neiva, 1912) (Diptera: Psychodidae: Phlebotominae) in Campo Grande, an endemic area of visceral leishmaniasis in Mato Grosso do Sul, Brazil,” Acta Tropic, vol. 107, no. 3, pp. 238–241, 2008.
[6] A. M. Marassá, C. A. Consales, and E. A. B. Galati, “Enzyme-linked immunosorbent assay biotin/avidin method standardization, for identification of Lutzomyia (Lutzomyia) longipalpis bloodmeals (Lutz & Neiva, 1912),” Revista da Sociedade Brasileira de Medicina Tropical, vol. 37, no. 6, pp. 441–446, 2004.
[7] W. D. Sudia and R. W. Chamberlain, “Battery-operated light trap, an improved model,” Mosquito News, vol. 2, pp. 126–129, 1962.
[8] H. Pugedo, R. A. Barata, J. C. França-Silva, J. C. Silva, and E. S. Dias, “HP: an improved model of suction light trap for the capture of small insects,” Revista da Sociedade Brasileira de Medicina Tropical, vol. 38, no. 1, pp. 70–72, 2005.
[9] T. R. Burクト, W. G. Goodman, and G. R. DeFoliart, “Identification of mosquito blood meals by enzyme-linked immunosorbent assay,” American Journal of Tropical Medicine and Hygiene, vol. 30, no. 6, pp. 1336–1341, 1981.
[10] R. Duarte, “Ensaios imunoenzimáticos ELISA para identificação experimental de fontes alimentares em Panstrongylus megistus (Burmeister, 1835) (Hemiptera: Reduviidae),” Tese de Mestrado, Instituto Oswaldo Cruz/FIOCRUZ, 1997.
[11] D. C. Young and N. A. Duncan, “Guide to the identification and geographic distribution of Lutzomyia sand flies in México, the West Indies, Central and South America (Diptera: Psychodidae),” Memoirs of the American Entomological Institute, vol. 54, pp. 1–881, 1994.
[12] L. M. Deane and Leishmaniose visceral no Brasil, “Estudos sobre reservatórios e transmissores realizados no Estado do Ceará. Rio de Janeiro, Serviço Nacional de Educação Sanitária,” Rio de Janeiro, pp.162, 1962.
[13] R. Rosabal and A. Trejos, “Phlebotomus de El Salvador (Diptera: Psychodidae). II. Observaciones sobre su biologia con especial referencia a P. longipalpis,” Revista de Biología Tropical, vol. 13, pp. 219–228, 1965.
[14] R. Zeledon, J. Murillo, and H. Gutierrez, “Ecology of Lutzomyia longipalpis (Lutz & Neiva, 1912) and possibilities of the existence of visceral leishmaniasis in Costa Rica,” Memorias do Instituto Oswaldo Cruz, vol. 79, no. 4, pp. 455–459, 1984.
[15] J. C. Araújo, J. M. M. Rebelo, M. L. Carvalho, and V. L. L. Barros, “Composição dos flebotomíneos (Diptera, Psychodidae) do Município de Raposo-MA, Brasil. Área endêmica de leishmanioses,” Entomol Vectores, vol. 7, pp. 33–47, 2000.
[16] M. L. Carvalho, J. M. M. Rebelo, J. C. Araújo, and V. L. L. Barros, “Aspectos ecológicos e parasitológicos de Leishmania (Diptera, Psychodidae) do Município de São José do Ribamar, MA, Brasil. Área endêmica de leishmanioses,” Entomol Vectores, vol. 7, pp. 19–32, 2000.
[17] F. O. Dias, E. S. Lorosa, and J. M. Rebelo, “Blood feeding sources and peridomiciliation of Lutzomyia longipalpis (Lutz & Neiva, 1912) (Psychodidae, Phlebotominae),” Cadernos de Saúde Pública, vol. 19, no. 5, pp. 1373–1380, 2003.
[18] B. Alexander, R. Lopes de Carvalho, H. McCallum, and M. H. Pereira, "Role of the domestic chicken (Gallus gallus) in the epidemiology of urban visceral leishmaniasis in Brazil," *Emerging Infectious Diseases*, vol. 8, no. 12, pp. 1480–1485, 2002.

[19] R. Lainson and E. F. Rangel, "Lutzomyia longipalpis e a eco-epidemiologia da leishmaniose visceral americana (LVA) no Brasil. In: Rangel EF e Lainson R (edts), Flebotomineos do Brasil," *Rio de Janeiro, Fiocruz*, pp. 311–336, 2003.

[20] M. S. Fernández, O. D. Salomón, R. Cavia, A. A. Perez, S. A. Acardi, and J. D. Guccione, "*Lutzomyia longipalpis* spatial distribution and association with environmental variables in an urban focus of visceral leishmaniasis, Misiones, Argentina," *Acta Tropica*, vol. 114, no. 2, pp. 81–87, 2010.

[21] J. G. D. Silva, G. L. Werneck, M. D. S. P. Cruz, C. H. N. Costa, and I. L. De Mendonça, "Natural infection of *Lutzomyia longipalpis* by *Leishmania* sp. in Teresina, Piauí State, Brazil," *Cadernos de Saúde Pública*, vol. 23, no. 7, pp. 1715–1720, 2007.

[22] V. L. F. Camargo-Neves, L. A. C. Rodas, and A. C. Gomes, "Avaliação do hábito alimentar de *Lutzomyia longipalpis* no Estado de São Paulo," *Boletim Epidemiológico Paulista*, vol. 4, pp. 2–7, 2007.

[23] S. M. da Silva, P. F. B. Rabelo, N. D. F. Gontijo et al., "First report of infection of *Lutzomyia longipalpis* by *Leishmania* (Leishmania) infantum from a naturally infected cat of Brazil," *Veterinary Parasitology*, vol. 174, no. 1-2, pp. 150–154, 2010.

[24] A. C. Morrison, C. Ferro, and R. B. Tesh, "Host preferences of the sand fly *Lutzomyia longipalpis* at an endemic focus of American visceral leishmaniasis in Colombia," *American Journal of Tropical Medicine and Hygiene*, vol. 49, no. 1, pp. 68–75, 1993.

[25] I. A. Sherlock, J. C. Miranda, M. Sadigursky, and G. Grimaldi Júnior, "Natural infection of the opossum *Didelphis albiventris* (Marsupialia, Didelphidae) with *Leishmania donovani*, in Brazil," *Memorias do Instituto Oswaldo Cruz*, vol. 79, no. 4, p. 511, 1984.

[26] I. A. Sherlock, J. C. Miranda, M. Sadigursky, and G. Grimaldi Jr., "Experimental infection of the opossum *Didelphis albiventris* (Marsupialia, Didelphidae) with *Leishmania donovani*," *Memorias do Instituto Oswaldo Cruz*, vol. 83, no. 1, p. 141, 1988.

[27] H. D. F. H. Schallig, E. S. da Silva, W. F. van der Meide, G. J. Schoone, and C. M. F. Gontijo, "*Didelphis marsupialis* (Common Opossum): a potential reservoir host for zoonotic leishmaniasis in the metropolitan region of Belo Horizonte (Minas Gerais, Brazil)," *Vector-Borne and Zoonotic Diseases*, vol. 7, no. 3, pp. 387–393, 2007.

[28] M. A. A. Cabrera, A. A. Paula, L. A. B. Camacho et al., "Canine visceral Leishmaniasis in Barra de Guaratiba, Rio de Janeiro, Brazil: assessment of risk factors," *Revista do Instituto de Medicina Tropical de Sao Paulo*, vol. 45, no. 2, pp. 79–83, 2003.

[29] A. C. Gomes, "Vigilância Entomológica," *Informe Epidemiológico do SUS*, vol. 11, pp. 79–80, 2002.