

Inventory of potential vectors of trypanosoma and infection rate of the Tsetse fly in the National Park of Ivindo, Gabon.

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Abstract

Background: *Trypanosoma*'s vectors distribution is poorly investigated in Gabon, where Trypanosomiasis historical foci exist. Thus, an active detection of *Trypanosoma sp* transmission needs to be assessed.

Objectives: The present study aims to identify potential vectors of *Trypanosoma sp* and to evaluate the infection rate of the Tsetse fly in an area of Gabon.

Methods: An entomological survey was conducted in the National Park of Ivindo in May 2012 using Vavoua traps. All captured insects were identified. Tsetse were dissected and organs were microscopically observed to detect the presence of *Trypanosoma sp*.

Results: 247 biting flies known as vectors of Trypanosomiasis were caught including 189 tsetse flies, 32 *Tabanid* and 26 *Stomoxys*. Tsetse flies had the highest bulk densities per trap per day (ADT = 3 tsetse / trap / day), while the lowest density was found among *Stomoxys* (ADT= 0.41 *Stomoxys* / trap / day). The infection rate of flies was 6.3%. Infectious organs were midguts and to a lesser extent salivary glands and proboscis.

Conclusion: The presence of Tsetse infected by *Trypanosoma* highlights an existing risk of trypanosomiasis infection in the National Park of Ivindo.

Keywords: Tsetse fly, *Tabanids*, vector, *Stomoxys*, biodiversity, trypanosomiasis.

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Introduction

Human African Trypanosomiasis (HAT) or sleeping sickness threatens 70 million of people in sub-Saharan Africa¹. Without adequate treatment, sleeping sickness is a fatal disease. However, less than 10% of at-risk populations are monitored. Since 2009, less than 10000 of HAT cases have been reported, but the World Health Organization (WHO) estimates that the number of infected persons is near of 20000¹. HAT is endemic in 35 countries of the African region with variable levels and

is located in more than 250 active foci mainly in Angola, Democratic Republic of Congo and Sudan^{2,3}.

In Central Africa, trypanosomiasis is endemic in all ecological features (forest, savannah, mangrove, river valleys, islands, etc.)⁴. In Gabon, a historic landmark of the HAT in Central Africa, the disease was endemic in seven of the nine provinces during the last century⁵. Recently new cases from the provinces of the Estuaire and Ogooué Maritime have been reported (reports Programme National de Lutte contre la Trypanosomose Humaine Africaine). The Komo-Mondah, inside Estuaire, is the most active focus of HAT in Gabon with an average of 20 cases per year⁶. In this focus, *Trypanosoma sp*, transmitted by tsetse flies and mechanically by tabanids and stable flies, have been found⁷; notably *Trypanosoma brucei gambiense*, a human *Trypanosoma* and *Trypanosoma congolense* and *Trypanosoma vivax*, cattle parasites^{6,8}. Nevertheless, other existing old foci of the disease are not monitored since several years; thus the proportion of persons at risk of acquiring the disease is presumably underestimated in the country.

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In the region of Ogooué Ivindo, in the east of country, old foci were mainly located along rivers and tributaries⁹. Previous work on the abundance and distribution of blood sucking *Diptera* revealed a high density of stable flies, tabanids and tsetse flies in the Bai located near the National Park of Ivindo¹⁰. The aim of the study was to determine the density of *Trypanosoma* vectors and to assess the infection rate of the tsetse fly, in the National Park of Ivindo in Gabon, fifteen years after the last survey.

Methods:

Study site:

The study was conducted in May 2012 at Ipassa, situated in the NPI, in the region of the Ogooué-Ivindo, in the North-Eastern part of Gabon. It is a forest area located at 632 km from Libreville, the capital city of Gabon¹¹. The National Park of Ivindo has a wide biodiversity and its ecology is already described¹¹⁻¹². The climate is equatorial, characterized by an alternation of two dry (mid-December to mid-March and mid-June and mid-September) and two rainy seasons (mid-March to mid-June and mid-September to mid-December). The average annual rainfall is about 1600-1800 mm. The mean temperature is about 24 ° C with a minimum of 21.7 ° C in July and a maximum of 25 ° C in April. The vegetation is composed by primary, secondary forests, and flood plain forests. The fauna is rich and diverse¹². The main activities of the inhabitants are fishing, hunting and agriculture.

Entomological survey:

Sessions fly trapping were conducted during the long rainy season with Vavoua traps¹³. A network of nine traps was set up during seven days. All traps were placed in the forest environment along the Ivindo river, 500 m away from each other. Each trap site was cleared to ensure reasonable visibility of the trap and was activated before 7:00.am and withdrawn after 5:00.pm. When the traps are controlled, each capture cage, contained in the trap, is recovered, numbered and dated; and during the day sent to the laboratory, at the station of the Institute for Research in Tropical Ecology Research located at 12 km from the city of Makokou.

Fly identification and *Trypanosoma* detection in tsetse:

The fly's species identification was performed in the laboratory. Morphological discrimination between dif-

ferent fly's taxa (*Glossina*, *Stomoxys*, *Tabanids*) was made according to the species identification keys: the Oldroyd key for *Tabanidae* group, Pollock and Brunhes for tsetse flies group and the Zumpt key for the stable flies group (*Stomoxys*)^{14,15,16,17}. All observations and dissections were performed under a magnifying glass.

After sorting, identification and counting, tsetse flies were dissected (within 24h following the capture) into a drop of sterile 0.9% saline water. Midgut, proboscis and salivary glands were then examined with a light microscope at a 400x magnification for the detection of *Trypanosoma* in the trypomastigotes procyclic form. Teneral flies were identified by the residual bag of the larval stage in the midgut¹⁸. Organs, both infected and uninfected were transferred separately into a microfuge tubes containing 50µl of 70% ethanol and stored in -20°C for further molecular analysis.

Data analysis

Data were recorded and analyzed using Excel.

The abundance of each flies species was defined by the apparent density per trap and per day (ADT) as follows:

$$ADT = \frac{\text{Number of flies captured}}{\text{Number of traps} \times \text{number of days of capture}}$$

The sex-ratio was determined as follows: Number of males / Number of females

The fly infection rate was calculated using the following formula:

$$\text{Infection rate} = \left(\frac{\text{Number of infected flies}}{\text{Total Number of dissected flies}} \right) \times 100$$

The *Trypanosoma* risk index =(ADT x infection rates) was also calculated.

The *Margalef* diversity index was calculated to assess the diversity of the fly species in the study area. This index is calculated using the following formula: $D = (S-1) / \log N$, where "S" is the number of species and "N" the total number of individuals collected¹⁹.

Results:

A total of 247 biting flies, belonging to the following families *Tabanidae*, *Muscidae* and *Stomoxysidae*, were collected. They were distributed into 5 genus: *Glossina* (n=189), *Stomoxys* (n= 26), *Chrysops* (n= 23), *Tabanus* (n= 7) and *Ancala* (n= 2). The apparent density per trap and per day (ADT) was 3G/t/d for *Glossina*, 0.4S/t/d for *Stomoxys* and 0.5T/t/d for *Tabanid*.

Species identification:

Among tsetse flies, the predominant species was *Glossina palpalis palpalis* (n= 114, 60.3%) (Table 1). *G. fusca congolensis*, *G. nashi*, *G. tachinoïdes*, *G. tabaniformis* and *G. frezili* were also identified in the study site (table 1). In this group the Margalef index was 2.2. Teneral flies (n= 9) were collected in the biotope including *Glossina palpalis palpalis* (n= 4), *Glossina fusca congolensis* (n= 2), *Glossina nashi* (n= 2) and *Glossina frezili species* (n=1).

Among *Tabanidae* (n= 32), *Tabanus*, *Chrysops* and *Ancala genus* were determined.

The most abundant species identified was *Chrysops silaceus* (53%; n=17). *Chrysops dimidiata* (n=6), *Tabanus sp* (n=5), *Tabanus taeniola* (n=2), *Ancala nilotica* (n=1), *Ancalafasciata* (n=1) were also found. The Margalef index was of 3.3.

Out of the four *Stomoxys* species found during the survey, *Stomoxys inornatus* (73%; n=19) predominated although *Stomoxys nigra nigra* (n=3), *Stomoxys indica* (n=3) and *Stomoxys omega* (n=1) were also identified. *Stomoxys Margalef* index was of 2.1.

Table 1: abundance and infection rate of the tsetse fly

Tsetse flies			Gender		Infection rate	
	Species	N	%	Male, n	Female, n	Male, %(n)
<i>G. palpalis palpalis</i>	114	60.3	55	59	5.4 (3)	10.1 (6)
<i>G. fusca congolensis</i>	36	19.1	16	20	0.0 (0)	0.0 (0)
<i>G. nashi</i>	14	7.4	5	9	0.0 (0)	0.0 (0)
<i>G. tachnoïdes</i>	11	5.8	4	7	0.0 (0)	14.3 (1)
<i>G. tabaniformis</i>	7	3.7	3	4	33.3 (1)	25.0 (1)
<i>G. frezili</i>	7	3.7	5	2	0.0 (0)	0.0 (0)
Total	189	100	88	101	4.5 (4)	7.9 (8)

Sex-ratio and rate of infection among Tsetse flies:

The sex-ratio of the captured tsetse flies was 0.9 (Table 1). All tsetse flies were dissected. Among them, 12 were found infected out of which nine were *G. palpalis palpalis*, two *G. tabaniformis* and one *G. tachinoïdes*. The infection rate was of 6.6% (n=12/180) in the non teneral flies. The *Trypanosoma* risk index was of 0.19. The sex ratio of infected flies was 0.5. Females were more infected: 7.9% infected females (n=8/101) and 4.5% (n=4/88) infected males. The infection was detected in

the midgut of all flies. In five tsetse flies, salivary glands and proboscis were additionally infected.

Discussion:

The control and the monitoring of HAT transmission in historical focus where the disease is considered eliminated are based on selective serological surveillance². However, entomological surveys allowing the vectors and reservoir identification as well as the parasite detection are required to assess the local transmission of

Trypanosoma sp. In the present study, investigations on *Trypanosoma sp* vectors distribution in an old focus of the National Park of Ivindo in Gabon showed a high prevalence of Tsetse flies, vectors of HAT, as well as stable flies and *Tabanids* among circulating flies. *Stomoxys* and *Tabanids*, potentials vectors of *Trypanosoma sp*, represented 10% and 3% of the whole flies collected, respectively.

Overall, 15 species of flies have been identified, except five from the *Tabanidae* family. Within each group, between four and six species were identified; comparable Margalef index was found between tsetse flies and *Stomoxys*. In a previous study carried out in the area of Ogooué Ivindo, precisely along a transect in the Baï, 22 different species were detected during the rainy season; 14 *Tabanids*, five *Glossina* and three *Stomoxys* species²⁰. The location of the study may influence the fly's species diversity. NPI and the Baï, which is at a distance of 160 km, share the same fly species. Nevertheless, *G. frezili*, *G. nashi*, *S. indica*, *A. nilotica*, *A. fasciata* and *C. dimidiata* were not found in the Baï during the rainy season. In this forest of Ogooué Ivindo, all environmental components such as temperature (15°C– 25°C), humidity and light are found for the ubiquitous distribution of the tsetse fly²¹.

The group of Tsetse flies, the single biological vectors of trypanosomiasis, was the most abundant in the park. Females were more frequently captured. Similar observations were previously reported and were related to the technique of capture^{22,23}. It is suggested that *Glossina* females are more frequently captured by Vavoua trap, as observed in the present study²². Among the captured Tsetse flies, less than 5% were teneral, that is a good indicator of tsetse reproduction sites; while none was found in a study carried out in Ivory Coast out of 269 flies examined during the rainy and dry seasons²².

The present study is the first performed after more than fifteen years to investigate the *Glossina* infection rate in the region of Ogooué Ivindo⁶. Identification of infected tsetse flies reveals the existence of a local transmission cycle of *Trypanosoma sp* at the NPI. *Glossina palpalis palpalis*, the main vector of human Trypanosomiasis, was the main fly species infected, mostly females as reported elsewhere^{21,8,24,25}. The predominance of females among infected flies is probably due to the greater number of meals taken compared to male and to their

longer life²³. *Trypanosoma sp* was specifically detected in the midgut and secondarily (or not) in salivary glands, as reported by others^{26,27}.

The infection rate in the present study, although determined after microscopic examination, is higher than the rate found by PCR, a highly sensitive method, in a neighboring country, Equatorial Guinea, precisely at Campo Rio (3.2%), suggesting the detection of a likely higher prevalence of infected flies if molecular tools were used in the present survey²⁷. Nevertheless, it is lower compared to the rate reported in the suburban area of Libreville (9.3%) and in various areas from Ivory Coast ranging from 20 to 28%^{27,28}.

Between 2000 and 2009, 328 HAT cases have been reported in Gabon, among them, six were European cases whom one was exposed in a rural forest area^{3,29,30,31}. Moreover, data from epidemiological and entomological surveys carried out in Gabon, specifically, in the province of the Estuary and more recently, the Ogooué-Maritime region, showed the transmission of the disease in these areas^{5,6,24,25}. The present findings highlight the need of identification of species of *trypanosomes* endemic in the NPI and of the case detection by an active screening of the population at risk in the NPI and in the surrounding areas along the river.

Limitation

The main limitation of the study was the lack of *Trypanosoma* species molecular identification. Taking into account that the development of many insects may present some peak of abundances depending on climatic factors, the differences of abundance observed between the detected insect families should be taken with caution³².

Conclusion

Different species of blood-sucking flies, potential vectors of trypanosomiasis, have been identified at Ogooué Ivindo. The presence of infected tsetse flies in this locality, former HAT foci, and especially in the Ivindo reserve underlines the existence of a *Trypanosoma sp* transmission risk in this area. Although, a contact between tsetse flies and human must be demonstrated and the human *Trypanosoma sp* identified to confirm the local transmission of HAT, Ivindo is an old focus and the presence of infected flies highlight the importance of *Trypanosoma* species identification in this area.

Competing interests

The authors declare that they have no competing interests.

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