

## THE PREVALENCE OF AFRICAN ANIMAL TRYPANOSOMOSIS AND TSETSE PRESENCE IN WESTERN SENEGAL

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### Summary:

In 2005, the Government of Senegal initiated a tsetse eradication campaign in the Niayes and La Petite Côte aiming at the removal of African Animal Trypanosomosis (AAT), which is one of the main constraints to the development of more effective cattle production systems. The target area has particular meteorological and ecological characteristics that provide great potential for animal production, but it is unfortunately still infested by the riverine tsetse species *Glossina palpalis gambiensis* Vanderplank (Diptera: Glossinidae). The tsetse project in Senegal has adopted an area-wide integrated pest management (AW-IPM) approach that targets the entire tsetse population within a delimited area. During the first phase of the programme, a feasibility study was conducted that included the collection of entomological, veterinary, population genetics, environmental and socio-economic baseline data. This paper presents the parasitological and serological prevalence data of AAT in cattle residing inside and outside the tsetse-infested areas of the target zone prior to the control effort. At the herd level, a mean parasitological prevalence of 2.4 % was observed, whereas a serological prevalence of 28.7 %, 4.4 %, and 0.3 % was obtained for *Trypanosoma vivax*, *T. congolense* and *T. brucei brucei*, respectively. The observed infection risk was 3 times higher for *T. congolense* and *T. vivax* in the tsetse-infested than in the assumed tsetse-free areas. Moreover, AAT prevalence decreased significantly with distance from the nearest tsetse captured which indicated that cyclical transmission of the parasites by tsetse was predominant over mechanical transmission by numerous other biting flies present. The importance of these results for the development of a control strategy for the planned AW-IPM campaign is discussed.

**KEY WORDS:** African animal trypanosomosis, tsetse, cyclical transmission, mechanical transmission, area-wide integrated pest management, feasibility study, Senegal.

### Résumé : PRÉVALENCE DES TRYPANOSOMOSIS ANIMALES AFRICAINES ET PRÉSENCE DES GLOSSINES DANS L'OUEST DU SÉNÉGAL

En 2005, le Gouvernement du Sénégal a initié une campagne d'élimination des glossines dans la zone des Niayes et La Petite Côte dans le but de supprimer les Trypanosomoses Animales Africaines (TAA), qui font partie des contraintes principales au développement des systèmes de production intensifs. La zone cible présente des conditions climatiques et écologiques particulières avec un potentiel de production animale important, mais elle est malheureusement toujours infestée par l'espèce de glossine riveraine *Glossina palpalis gambiensis* Vanderplank (Diptera : Glossinidae). Le projet d'élimination des glossines a adopté une stratégie de gestion intégrée des ravageurs à l'échelle d'une population totale dans une zone donnée. La première phase du programme correspond à une étude de faisabilité qui inclut la collecte de données de base entomologiques, vétérinaires, génétiques, environnementales et socio-économiques. Cet article présente les données de prévalence parasitologique et sérologique dans et hors de la zone infestée par les glossines avant le début de la lutte. La prévalence parasitologique moyenne intra-troupeau était de 2,4 %, alors que la prévalence sérologique était de 28,7 %, 4,4 % et 0,3 % pour *Trypanosoma vivax*, *T. congolense* et *T. brucei brucei* respectivement. Le risque d'infection observé était trois fois plus élevé dans la zone infestée par les glossines que dans la zone considérée comme non-infestée. De plus, la prévalence diminuait significativement avec la distance à la glossine capturée la plus proche, ce qui indique que la transmission cyclique des parasites domine, par rapport à la transmission mécanique par de nombreux autres insectes piqueurs présents. L'importance de ces résultats pour le développement de la stratégie de lutte intégrée est discutée.

**MOTS CLÉS :** trypanosomose animale africaine, tsé-tsé, transmission cyclique, transmission mécanique, gestion intégrée des ravageurs, étude de faisabilité, Sénégal.

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## INTRODUCTION

The Niayes area and La Petite Côte are located along the Atlantic coast of Senegal and include four administrative districts: Dakar, Thies, Louga and Saint-Louis. Particular meteorological and ecological characteristics of this area provide great potential for agricultural development in general and animal production (cattle, donkeys, horses, small ruminants, pigs and poultry) in particular. Most of these animals are however susceptible to the debilitating disease African Animal Trypanosomosis (AAT) (Mulligan, 1970) which is widespread in sub-Saharan Africa (Itard *et al.*, 2003). AAT is

indeed the main constraint to the development of more effective cattle production systems in African countries infested with tsetse (Itard *et al.*, 2003), their cyclical vectors. Details of the infestation of the Niayes area and Petite Côte with *Glossina palpalis gambiensis* Vanderplank (Diptera: Glossinidae) were already collected in the 1960s (Morel & Touré, 1967; Touré, 1971, 1974), which was followed by eradication attempts in the 1970s and 1980s using mainly ground spraying of residual insecticides (Touré, 1973). Thereafter, the tsetse and trypanosomiasis problem seemed to have disappeared, which allowed the introduction of exotic cattle breeds in the area (Ba Diao, 2005) but flies were detected again in 1998 (unpublished report of the Direction des Services Vétérinaires Services (DSV)). The maintenance of these exotic cattle (Holstein, Jersey, Montbeliard, etc) in zero ground-grazing units has only been possible by keeping these trypano-susceptible animals under constant prophylactic treatment with trypanocidal drugs, which is not only cost ineffective but not sustainable in the long term. AAT remains a major pathological problem in the area, where the mean daily milk production is still limited to 6.9 kg (s.d. 3) despite much higher genetic potential of these exotic breeds (Ba Diao, 2005). In 2005, the DSV initiated a campaign called «Projet de lutte contre les glossines dans les Niayes» with the goal to develop a sustainable solution to the tsetse and trypanosomiasis problem in the Niayes. The programme is funded by the Government of Senegal and technically and financially supported by the International Atomic Energy Agency (IAEA), the Food and Agriculture Organization of the United Nations (FAO) and the Centre de Coopération Internationale en Recherche Agronomique pour le Développement (CIRAD). The project is being implemented in the context of the Pan African Tsetse and Trypanosomiasis Eradication Campaign (PATTEC), a political initiative of the African heads of state that calls for increased efforts to manage the tsetse and trypanosomiasis problem in sub-Saharan Africa ([http://www.africa-union.org/Structure\\_of\\_the\\_Commission/depPattec.htm](http://www.africa-union.org/Structure_of_the_Commission/depPattec.htm)). The tsetse project in Senegal has adopted an area-wide integrated pest management (AW-IPM) approach that aims at integrating various control tactics (e.g. traps and insecticide-impregnated targets (Bauer *et al.*, 2005; Green, 1994), live baits (Bauer *et al.*, 1995; Bouyer *et al.*, 2009; Bouyer *et al.*, 2007), and the sterile insect technique (SIT) (Dyck *et al.*, 2005)) to target an entire tsetse population within a circumscribed area (Klassen, 2005).

As a first phase of this project, entomological, veterinary, and population genetics baseline data have been collected as part of a feasibility study (Bouyer *et al.*, 2010; Solano *et al.*, 2010). The collection of socio-economic and environmental data will complete the feasibility study effort. The DSV, the Institut Sénégalais de

Recherches Agricoles (ISRA), and the Centre de Suivi Ecologique (CSE) implemented this feasibility study to assess whether a sustainable zone free of *G. p. gambiensis* can be created in the Niayes and La Petite Côte. In addition, the data collected would constitute a base of reference data that would allow the evaluation of the progress and cost-effectiveness of a future control campaign. For the entomological baseline data study, a stratified sampling procedure was developed using geographical information systems (GIS), remote sensing data and probability models to ascertain the current distribution of tsetse in the target area (Bouyer *et al.*, 2010). Trapping data from 683 unbaited Vavoua traps deployed between January 2008 and February 2009 showed that *G. p. gambiensis* was the only species present in the area and was distributed in an area of 525km<sup>2</sup> of the Niayes and La Petite Côte (Fig. 1). No tsetse flies were sampled between the Niayes and the Sine Saloum (Missira), which is situated more than 100 km to the south-east of the Niayes. The zero catches in this area provided the first indications that the *G. p. gambiensis* populations of the Niayes were isolated from the remainder of the tsetse belt in south-eastern Senegal (Bouyer *et al.*, 2010). Using three markers (microsatellite DNA, mitochondrial COI DNA, and geometric morphometrics) a population genetic study revealed the absence of gene exchange between the *G. p. gambiensis* populations in the Niayes and those of the Sine Saloum (Missira), confirming the results of the entomological surveys that the populations of the Niayes can be considered isolated (Solano *et al.*, 2010). This paper presents the results of the veterinary survey, i.e. the parasitological and serological prevalence data of AAT in cattle in the target zone prior to the control effort, both inside and outside the tsetse-infested areas.

## MATERIELS AND METHODS

### STUDY AREA AND PERIOD

Between 30 August and 26 October 2007, the veterinary survey was conducted in 38 sites, which were geo-referenced using global positioning system (GPS) units (Fig. 1). These sites were randomly selected, since the tsetse distribution was not known at that time. In each of the sampling sites, blood was taken from the jugular vein of cattle that were declared sedentary to the Niayes and La Petite Côte by their owners. In herds that contained more than 50 animals, sampling was limited to 50 animals, whereas in smaller herds, all animals were sampled. From each animal, one blood sample was taken in a dry vacutainer tube and another in a vacutainer tube containing the anticoagulant ethylene diamine tetra acetic acid (EDTA). Blood samples were collected from in total 1,332 cattle belonging to 74 herds.

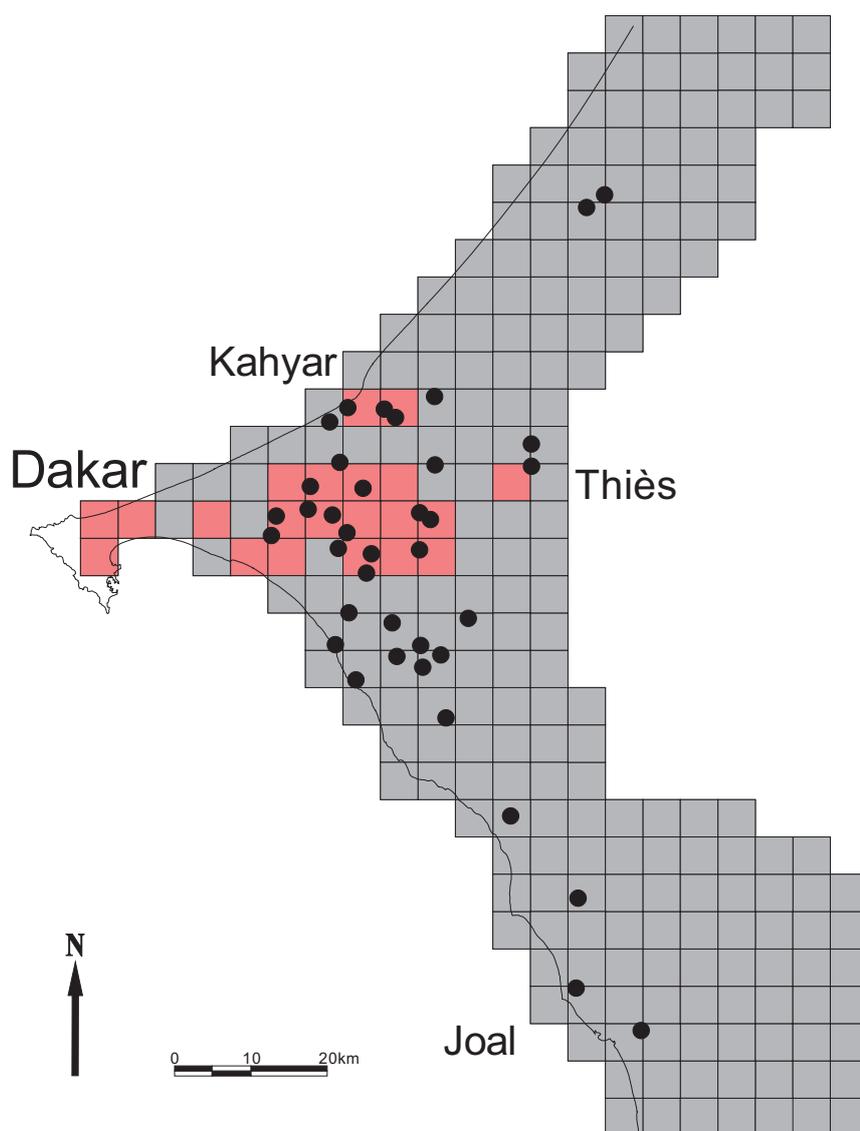


Fig. 1. – Location of the sampled herds and distribution of *Glossina palpalis gambiensis* in the Niayes area and La Petite Côte (presence = red, absence of record = grey).

#### LABORATORY ANALYSES

Three diagnostic techniques were used to assess AAT prevalence: direct examination of the Buffy coat using the dark ground hematocrite centrifuge technique (DG-HCT) for morphological identification (G\*20), a serological analysis using the Ab-enzyme-linked immunosorbent assay (ELISA) (Desquesnes *et al.*, 2001) and Polymerase Chain Reaction (PCR) (Desquesnes & Davila, 2002; Lefrançois *et al.*, 1998) on the positive samples to confirm the species of trypanosome involved. For the DG-HCT, blood subsamples were centrifuged in hematocrite tubes at 15,000 rounds/min during 5 minutes. The interphase of 1,141 samples was examined between slide and thin strip using a phase contrast microscope. All the 1,332 serum samples were tested for *Trypanosoma brucei brucei*, *T. congolense* and *T. vivax* using the Ab-ELISA technique. Monospecific primers of *T. vivax*, *T. brucei* sensu lato and *T. congolense* savannah

type (Desquesnes & Davila, 2002; Lefrançois *et al.*, 1998) were used for a PCR analysis which was carried out at the Centre International de Recherche-Développement sur l'Élevage en zone Sub-humide (CIRDES) in Burkina Faso.

#### STATISTICAL ANALYSES

The relationship between AAT prevalence and (1) tsetse presence, (2) tsetse apparent density, and (3) distance to the nearest trapped tsetse fly was assessed. For the first analysis, the AAT prevalence of herds located inside and in a buffer of 5 km around the known tsetse-infested area (to account for herd grazing distance), henceforth denoted 'tsetse-infested area', was compared to that of the area where the probability of presence of tsetse was below 0.05, henceforth denoted 'assumed tsetse-free area' (Bouyer *et al.*, 2010). The normality of the distribution in each area was tested

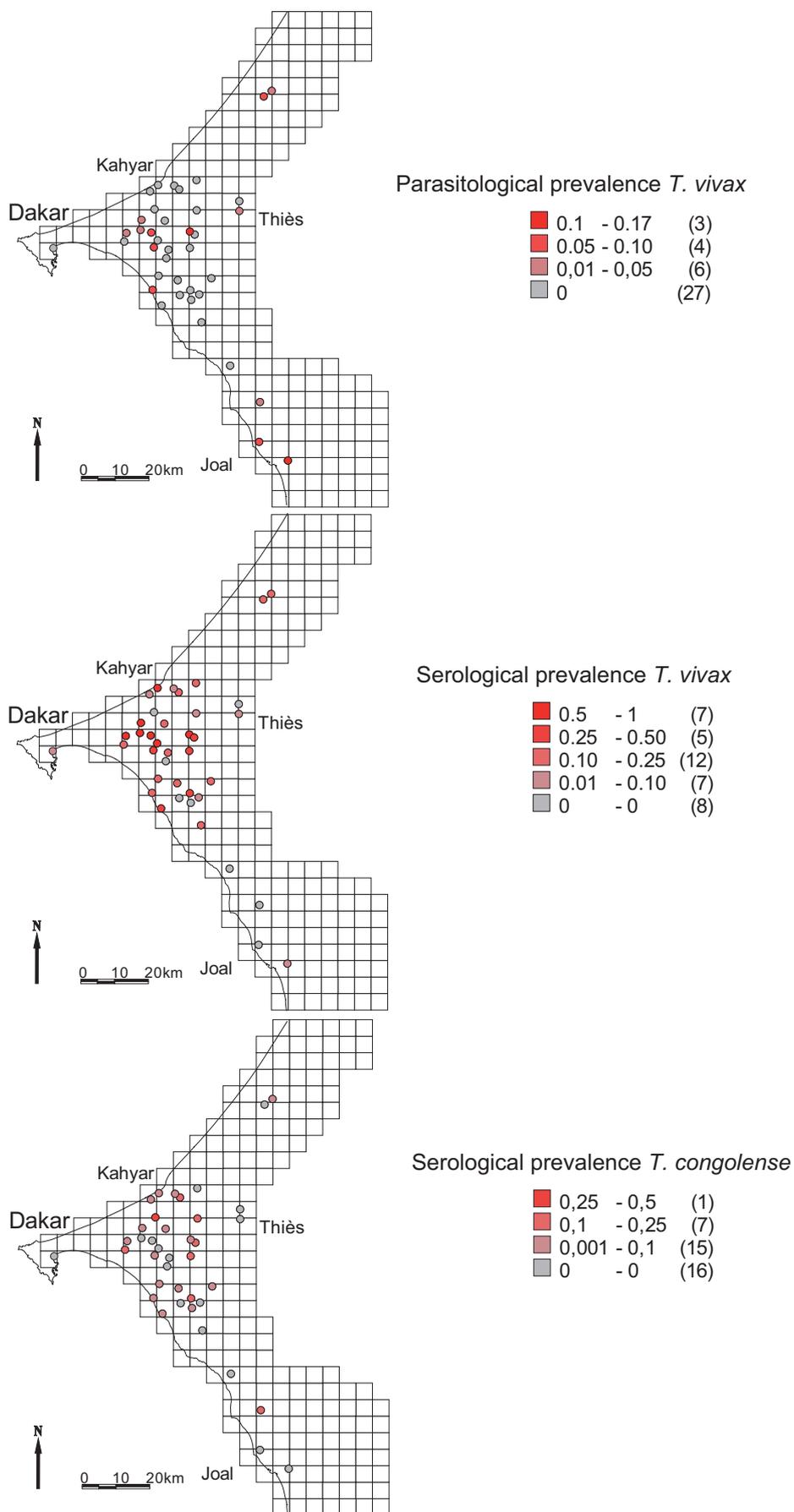


Fig. 2. – From top to bottom: herd parasitological prevalence of *T. vivax*; herd serological prevalence of *T. vivax* and *T. congolense* in the Niayes and La Petite Côte.

using the Kolmogorov-Smirnov test (Conover, 1971). Their variances were then compared using a F test and their means using a Student *t* test. To reduce the effects of animal mobility, the same analyses were repeated using only young cattle (< 3 years old) corresponding to 304 animals. Even if the herds are currently sedentary, old animals may still contain antibodies resulting from immune responses of past contacts with trypanosomes in tsetse-infested areas.

The correlation between the apparent density of tsetse and the AAT prevalence within the tsetse-infested area was assessed at the grid cell level [the area was subdivided in grid cells of 5x5 km to facilitate sampling procedures (Bouyer *et al.*, 2010)] using the Kendall's correlation test (Fig. 1).

Finally, the correlation between herd prevalence in young cattle and the distance to the nearest tsetse captured was investigated using Pearson's correlation test, after logarithmic transformation ( $\ln(x+1)$ ) of the herd prevalence rates.

## RESULTS

### PARASITOLOGICAL PREVALENCE

Animals positive for AAT were detected in 13 out of the 38 sites sampled. Among the 1,141 cattle analysed using the DG-HCT, 23 were positive, corresponding to a mean herd prevalence of 2.4 % (s.d. 4.3 %). Among the herds found positive with the DG-HCT, the prevalence varied between 2.6 % in Séguile Coopé (Mboro) and 13.3 % in Fadial (Joal) and Ndiakhaté (Pout) (Fig. 2). The parasitological analysis did not reveal the presence of *T. congolense* and all infections were attributed to *T. vivax*. No significant difference was observed between the parasitological prevalence of *T. vivax* in tsetse-infested and in the assumed tsetse-free areas ( $p = 0.88$ ). However, the parasitological prevalence of *T. vivax* in young animals was negatively correlated with distance to the nearest tsetse captured (Fig. 3,  $r = -0.41$ ,  $p = 0.016$ ). Inside the tsetse-infested area, the apparent density of tsetse was not correlated with the parasitological prevalence of *T. vivax* ( $p = 0.49$ ).

Among the 23 samples positive using the DG-HCT, a PCR analysis identified eight samples as *T. vivax*, whereas the others were negative for all trypanosome species. This might be due to a bad conservation of the samples during their transport from Senegal to Burkina Faso or to different *T. vivax* like trypanosomes.

### SEROLOGICAL ANALYSES

The overall mean serological prevalence for all samples was 28.7 % (s.d. 45.1 %), 4.4 % (s.d. 20.2 %), and 0.3 % (s.d. 4.7 %) for *T. vivax*, *T. congolense* and *T. brucei*

respectively. Fig. 2 shows the serological prevalence per herd for *T. vivax* and *T. congolense*.

The serological prevalence of *T. vivax* was significantly higher in the tsetse-infested (35.3 %, s.d. 33.5 %) than in the assumed tsetse-free area (13.9 %, s.d. 11.7 %) ( $p = 0.011$ ), with a difference of 21.5 % [95 % CI: 5.4-37.5 %]. The serological prevalence of *T. congolense* was not significantly different between the two areas ( $p = 0.09$ ).

This difference however became significant for both *Trypanosoma* species when only animals younger than three years were taken into consideration. In this age group the serological prevalence of *T. vivax* was 30.0 % (s.d. 33.6 %) in the tsetse-infested area and 8.3 % (s.d. 13.2 %) in the assumed tsetse-free area, corresponding to a difference of 21.7 % [95 % CI: 4.5-38.9 %] ( $p = 0.013$ ). For *T. congolense* a serological prevalence of 6.0 % (s.d. 9.7 %) was obtained for the tsetse-infested areas versus 1.2 % (s.d. 2.4 %) for the assumed tsetse-free areas ( $p = 0.05$ ), corresponding to a difference of 4.9 % [95 % CI: 0.3-9.4 %]. For this age group, the serological prevalence of *T. vivax* was also negatively correlated with distance to the nearest tsetse captured ( $r = -0.41$ ,  $p = 0.015$ ). This was not the case for *T. congolense* ( $p = 0.34$ ).

In the tsetse-infested area, the serological prevalence of *T. vivax* was highly correlated with the apparent density of tsetse ( $\tau = 0.423$ ,  $p = 0.008$ ). This correlation was not significant for *T. congolense* ( $p = 0.36$ ). The highest serological prevalence of *T. vivax* (96 %) and *T. congolense* (30 %) was observed in Diacksao Peuhl and Bahyakh, respectively, which are situated in the middle of the known tsetse-infested area.

## DISCUSSION

The Government of Senegal has recently embarked on a national program to intensify dairy production in the Niayes through the improvement of local cattle breeds through artificial inseminations with sperm from exotic, more productive breeds. Despite two tsetse control campaigns in the Niayes in the 1970s and 1980s (Touré, 1973), the tsetse fly *G. p. gambiense* and the disease AAT persisted and remain a serious impediment to the development of the livestock sector. The Ministry of Livestock and its partners (ISRA-LNERV, CSE, CIRAD, FAO/IAEA) have therefore initiated a study to assess whether the removal of the *G. p. gambiense* population from the Niayes using an area-wide integrated pest management approach (Vreysen *et al.*, 2007) will be feasible. Such a study would provide comprehensive data sets to serve as the underlying basis for the development of a scientifically sound intervention strategy and the necessary baseline data to monitor programme progress.

The results of the parasitological and serological survey reported in this paper indicated that *T. vivax* was the predominant trypanosome species in the Niayes with zero parasitological prevalence rates for *T. congolense*. These results are in accordance with observations in other sites of the northern tsetse belt in West Africa (Bouyer *et al.*, 2006; Guerrini & Bouyer, 2007) and might be due to a lower intrinsic vectorial capacity of *G. palpalis gambiensis* for *T. congolense* in comparison with e.g. *Glossina morsitans submorsitans* Newstead or *Glossina tachinoïdes* Westwood (Reifenberg *et al.*, 2008), two species common in West Africa but absent in the study area. Another hypothesis is a possible shorter lifespan of the *G. p. gambiensis* populations inhabiting the ecological marginal areas at the northern limit of their distribution, favouring trypanosome species with short extrinsic cycles (duration of the development of the parasite in the vector) (Bouyer *et al.*, 2006; Van den Bossche *et al.*, 2010).

The parasitological prevalence of *T. vivax* was low (2 %) but was probably under-estimated, since recent results obtained in The Gambia suggest that the sensitivity of the buffy coat technique is only 50 % (Faye *et al.*, 2001). Another factor that might have reduced the apparent parasitological prevalence is the constant prophylactic treatment of the animals with trypanocidal drugs. In addition, in many parts of the study area, *G. p. gambiensis* was mainly encountered in mango and other fruit orchards where cattle is prohibited and the flies mostly feed on humans and dogs. This likewise reduces

the transmission risk to cattle, lowers the prevalence rate and could also explain the high variability of the prevalence in herds located close to tsetse i.e. less than 5 km (Fig. 3).

The serological prevalence of *T. congolense* and *T. vivax* in young cattle was three times higher in the tsetse-infested than in the assumed tsetse-free areas. The presence of *T. congolense* in the Niayes is reported here for the first time and its wide distribution is worrying (Fig. 2) as the parasite is in general more pathogenic for cattle than *T. vivax* (Itard & Frézil, 2003).

The serological prevalence of *T. brucei* was too low to confirm its presence in the study area, given the negative parasitological results and the low specificity of ELISA test at the individual level (Desquesnes *et al.*, 2001). However, in December 2009, the presence of *T. brucei* s.l. was proven by a parasitological isolation from a wild *G. p. gambiensis* fly captured in Pout and fed on a goat.

The serological prevalence of *T. vivax* was positively correlated with tsetse density and negatively correlated with the distance to the nearest tsetse captured. The absence of a correlation between tsetse presence and density, the parasitological prevalence of both parasites and the serological prevalence of *T. congolense* is probably related to the very low values of the latter three parameters. Moreover, apparent fly density is not a good risk indicator and should be multiplied by tsetse infectious rate to better represent cyclical transmission (Bouyer *et al.*, 2006; Guerrini & Bouyer, 2007). However,

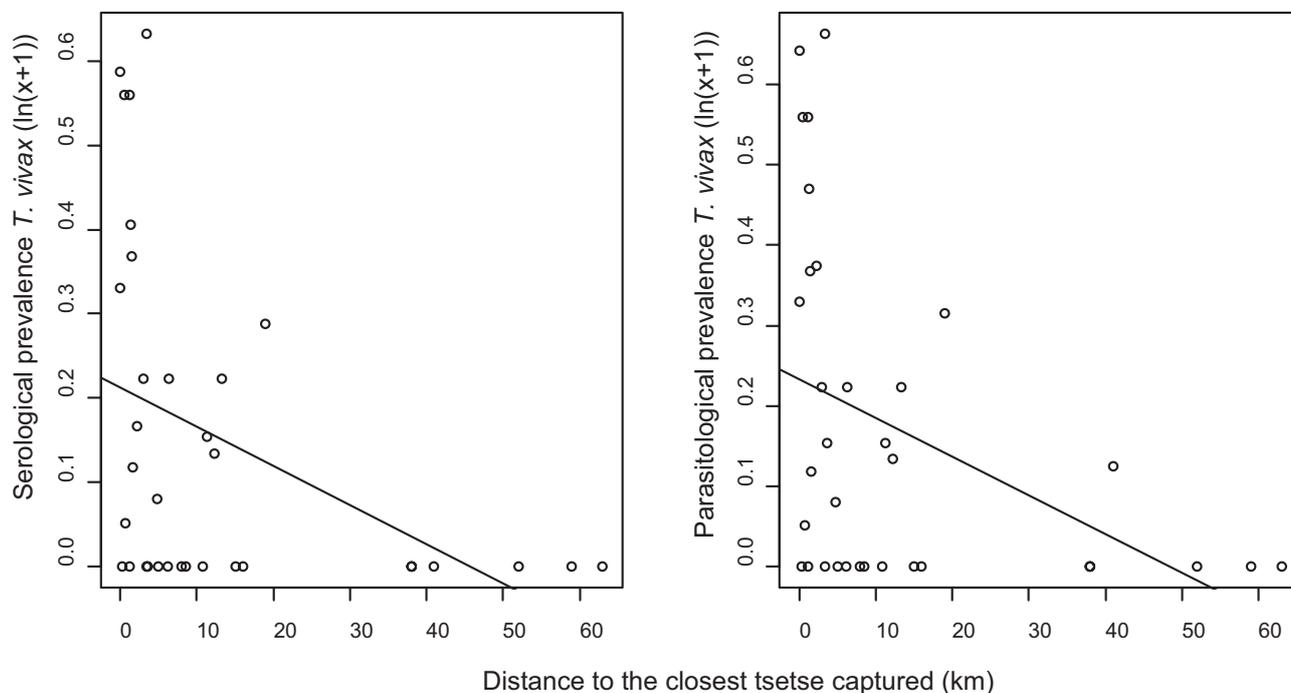


Fig. 3. – Correlation between serological (left) and parasitological (right) prevalence of *T. vivax* ( $\ln(x+1)$ ) and the distance to the nearest tsetse (*G. p. gambiensis*) captured.

the importance of cyclical transmission of AAT in the study area is obvious from the presented results, even if evidence exists that *T. congolense* and *T. vivax* can also be transmitted mechanically by biting flies (Desquesnes *et al.*, 2009; Desquesnes & Dia, 2003a, b, 2004). It is therefore likely that the sustainable removal of the *G. p. gambiensis* population from the Niayes and La Petite Côte will greatly reduce the prevalence of AAT, possibly followed by its disappearance as was demonstrated on Unguja Island. Both *T. congolense* (which was the dominant species before the start of the eradication campaign) and *T. vivax* disappeared from the island following the eradication of *Glossina austeni* Newstead (Vreysen *et al.*, 2000) despite very high densities of biting flies such as Stomoxyinae (Saleh *et al.*, 1999). These biting flies were therefore not capable to maintain the transmission cycle of the trypanosomes after the removal of the cyclical vector, raising questions on the epidemiological importance of biting flies. Similar observations were made in Sidéradougou, Burkina Faso after the removal of *G. p. gambiensis*, *G. tachinoides* and *G. morsitans submorsitans* from 3,500 km<sup>2</sup> of agropastoral land (Cuisance *et al.*, 1984; Delafosse *et al.*, 1996; Politzar & Cuisance, 1984). It would therefore be very useful to assess the epidemiological dynamics and the potential role of these biting flies after the completion of the tsetse eradication campaign in the Niayes and La Petite Côte.

During the entomological surveys, many species of Tabanidae and Stomoxyinae were captured throughout the target area, but at low mean apparent densities of 0.2 flies/trap/day (s.d. 0.6) and 1.9 flies/trap/day (s.d. 12.4) respectively. In certain areas, apparent densities were as high as 5.7 flies/trap/day for Tabanidae and 286 flies/trap/day for Stomoxyinae. The serological prevalence of both parasite species was not zero in the assumed tsetse-free areas, indicating a possible role of these biting flies outside the tsetse area. However, the mobility of cattle and their dispersal in and out of the tsetse-infested area should not be neglected and might have resulted in serological positive animals in the assumed tsetse-free areas. Moreover, it is possible that small tsetse pockets were not detected despite intensive trapping efforts inside the assumed tsetse-free area. From the models available, the most likely scenario in case of tsetse clearing is a disappearance of *T. congolense* (the most pathogen species for cattle) but it is possible that *T. vivax* persist with a different epidemiological pattern, i.e. periodical epidemics instead of an endemic transmission (Desquesnes *et al.*, 2009).

The severity of the symptoms caused by AAT is highly related to the type of cattle breed present in the various regions (Dayo *et al.*, 2009; Toure, 1997). In Basse Casamance in south-eastern Senegal, AAT seem less virulent due to the predominance of the trypanotolerant Ndama

cattle. In the areas with zebu cross-breeds, which are mostly located north to the tsetse belt, AAT is considered more virulent as compared to the Niayes (Touré, 1972). The serological and parasitological prevalence measured in this study are similar to the results of previous studies carried out in 1965 and 1969, which also revealed high variations in the prevalence rates between the various localities of the Niayes and La Petite Côte. In places where the forests have been completely replaced with agricultural areas, cattle are kept far from the tsetse-infested habitats and the prevalence remains low. The prevalence was however much higher in those areas where the preferred tsetse habitats have been preserved (e.g. Sangalkam, Noflaye, and Bambilor) and where parasitological prevalences of *T. vivax* up to 50 % have been observed during the rainy season (Touré, 1972). In southern Senegal, *T. congolense* predominates (65 % of the infections) (Fall *et al.*, 1993) and even in trypano-tolerant breeds such as Ndama, AAT reduces significantly animal draught power and hematocrite (Seck *et al.*, 2002).

The parasitological survey complemented the entomological baseline data collection effort and aimed at the indirect detection of presence of tsetse (Barclay & Hargrove, 2005; Vreysen, 2005). As a result, in sites with a herd prevalence above 10 % and where the distance to the nearest tsetse fly captured was more than 10 km (e.g. Mboro, Joal), additional entomological sampling was carried out. Despite these intensified trapping efforts of up to 30 days, no additional tsetse populations were detected. The development of a sound control strategy that aims at eradication of the *G. p. gambiensis* population and the removal of AAT in the Niayes and La Petite Côte will need to take into account both the data emanating from the entomological and the described parasitological and serological surveys. Even when further trapping failed to detect tsetse, the presence of parasitological and serological positive animals could be an indication of the presence of undetected relic pockets of tsetse flies and these zones should be considered as part of the target area of the area-wide eradication effort. The data of the surveys presented in this paper corroborate the outcome of the entomological data collection (Bouyer *et al.*, 2010) and the population genetics study (Solano *et al.*, 2010) that the *G. p. gambiensis* populations of the Niayes and La Petite Côte are isolated from the remainder of the tsetse belt in south-eastern Senegal.

The entomological, parasitological and serological surveys will be followed by a socio-economic study to evaluate the economical impact of AAT in this area, and to assess potential benefits of its elimination in relation to the total cost of the campaign. These data will not only be crucial to raise the necessary funds to implement the eradication campaign, but it will undoubtedly show

the great benefits that could emanate from an AW-IPM campaign aimed at the creation of a sustainable free zone of *G. p. gambiensis* in Western Senegal.

## ACKNOWLEDGEMENTS

We are very thankful to the veterinary staff involved in the field studies, particularly Abdou Gaye Mbaye, Babacar Camara, Babel Sow, Aïda Gueye Thiam, Mamadou Demba, Alphonse Manga, Mahanta Ndao, Alioune Ndiaye and Mansour Fall. We are indebted to the Direction des Services Vétérinaires and ISRA-LNERV for providing excellent working conditions.

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Reçu le 5 mai 2010

Accepté le 19 mai 2010