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## Nutritional stress of adult female tsetse flies (Diptera: Glossinidae) affects the susceptibility of their offspring to trypanosomal infections

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### ABSTRACT

The epidemiology of tsetse-transmitted trypanosomiasis depends, among other factors, on the proportion of infected flies in a tsetse population. A wide range of intrinsic and extrinsic factors seem to determine the ability of a tsetse fly to become infected and to transmit the parasite. In this paper, we investigated the effect of nutritional stress of reproducing female *Glossina morsitans morsitans* on the susceptibility of their offspring to trypanosomal infections. Adult female flies that were nutritionally stressed by feeding only once a week, produced pupae with a significant lower weight and offspring with a significant lower fat content as well as a lower baseline immune peptide gene expression. Moreover, infection experiments showed that the emerging teneral flies were significantly more susceptible to a *Trypanosoma congolense* or *Trypanosoma brucei brucei* infection than flies emerging from non-starved adult females. These findings suggest that in the field, substantial nutritional stress of adult tsetse flies, as is often experienced during the hot dry season, can increase significantly the vectorial capacity of the emerging teneral flies and thus result in an increased infection rate of the tsetse population.

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### 1. Introduction

In Africa, tsetse-transmitted trypanosomiasis poses a serious threat to animals and humans. About 10 million km<sup>2</sup> of sub-Saharan Africa, extending over 37 countries, are infected by tsetse flies. The epidemiology of this important disease is determined largely by the proportion of infected tsetse flies. A range of intrinsic and extrinsic factors among which environmental conditions, have been identified to affect the tsetse's susceptibility to trypanosomal infections and, thus affect the overall infection rate of the tsetse population (Leak, 1998; Aksoy et al., 2003; Macleod et al., 2007). For example, infection rates in tsetse flies are positively correlated with ambient temperature to which puparia or adult flies are exposed (Taylor, 1932; Burt, 1946; Fairbairn and Watson, 1955; Ndegwa et al., 1992). Moreover, high ambient temperatures shorten the duration of the development cycle within tsetse (Fairbairn and Culwick, 1950). The underlying mechanisms for the observed effect of ambient temperature on the susceptibility to infection are not well understood. The

nutritional status of the tsetse fly at the time of the infective blood-meal also affects its susceptibility to infections with *Trypanosoma congolense* or *Trypanosoma brucei brucei*. Indeed, a period of starvation (4 days for teneral flies and 7 days for adult flies) lowers the developmental barrier for a trypanosome infection, especially at the midgut level of the fly (Kubi et al., 2006). Here, it has been hypothesized that the reduction in the fat body reserve as a result of starvation reduces the fly's ability to mount a competent immune response against the invading trypanosomes. It has been suggested that the tsetse's innate immune response affects the establishment and maturation of trypanosomes in the vector (Hao et al., 2001; Boulanger et al., 2002; Lehane et al., 2004; Attardo et al., 2006). This immune response is determined by a range of antimicrobial peptides among which attacin, defensin, cecropin and dipterin that are mainly synthesized in the fat body in response to a 'foreign' micro-organism. Hence, all factors that reduce the fat body level of a teneral or adult fly may result in concomitant changes in the fly's vector competence. In this paper, we investigated the effect of nutritional stress of reproducing tsetse females, on the susceptibility of their offspring to trypanosomal infection. Results of this study may explain how changes in the tsetse's environment, that cause considerable stress, may have repercussions on the epidemiology of tsetse-transmitted human or animal trypanosomiasis.

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## 2. Materials and methods

### 2.1. Tsetse flies

Tsetse flies *Glossina morsitans morsitans* Westwood from the colony maintained at the Institute of Tropical Medicine (Antwerp, Belgium) were used in the experiments. The origin of this tsetse colony and rearing conditions were described by [Elsen et al. \(1993\)](#). Flies from this colony were fed 4 times a week on healthy rabbits (=non-starved colony). In addition to the main colony, a second colony of 2220 adult reproducing female flies was established and nutritionally stressed by feeding the flies only once a week on healthy rabbits (=starved colony) for 9 weeks. Pupae produced by the female flies of both colonies were collected daily and weighed.

### 2.2. Trypanosome infection of tsetse flies

The clonal strains of *T. congolense* IL 1180 ([Geigy and Kauffman, 1973](#)) and *T. brucei brucei* AntAR1 ([Le Ray et al., 1977](#)) were used to infect the experimental flies. Series of 30 male flies (less than 32 h old) that emerged from pupae produced by females of the starved or non-starved colony were given a single bloodmeal on trypanosome infected anesthetized mice (NMR1). The mice were infected with either *T. congolense* or *T. b. brucei* showing a parasitaemia of  $10^{6.9}$  or  $10^{8.1}$  trypanosomes per ml, respectively. Mice were anesthetized with 0.05 ml of a 8/3 (v/v) mixture of Anesketin® (diluted to 5% ketamine in physiological water) and Rompun®. Only fully engorged flies were retained and fed afterwards on healthy rabbits three times per week. To avoid re-infection of the flies, these rabbits were replaced at weekly intervals.

### 2.3. Tsetse infection rate

Twenty-one days (for infections with *T. congolense*) or 30 days (for infections with *T. b. brucei*) after the infective bloodmeal, all surviving flies were dissected using the method described by [Llyod and Johnson, 1924](#). Microscopical examination of the midgut and mouthparts or midgut and salivary glands were conducted to determine the presence of *T. congolense* (infection of midgut and mouthparts) or *T. b. brucei* (infection of midgut and salivary glands), respectively.

The proportion of immature (procyclic) infections was calculated as the proportion of dissected flies that had a trypanosomal infection in the midgut whereas the proportion of mature (metacyclic) infections was calculated as the proportion of dissected flies infected in midgut and mouthparts (*T. congolense*) or midgut and salivary glands (*T. b. brucei*). The maturation rate was calculated as the proportion of midgut infection that developed into a mature infection.

### 2.4. Determination of the fat content in the offspring

Thirty (30) male flies that emerged from pupae of the starved or non-starved colony were killed immediately after emergence, their legs and wings were removed and the carcasses were dried at 80 °C to constant weight. Afterwards the dry weight of pools of three flies was determined. Fat body was extracted through chloroform extraction of lipids for three days with daily changes of chloroform ([Langley et al., 1990](#)). After fat extraction and drying to constant weight, the residual dry weight of the pools of three flies was determined. The fat content of the flies was calculated as the difference between the dry weight and the residual dry weight. Weighing was done using a Sartorius® (GMBH, A120S) analytical balance ( $10^{-4}$  g).

### 2.5. Immune peptide gene expression levels in the freshly emerged flies

To compare the baseline immune peptide gene expression level of male flies derived from pupae of the starved or non-starved adult females, total RNA was extracted from the whole abdomens of respectively 24 and 20 male flies. RNA extraction was conducted in 1 ml Tripure® reagent (Roche) per sample following the manufacturer's instructions. Afterwards, the mRNA in 400 ng of a total RNA extract was reverse transcribed using Oligo(dT)<sub>15</sub> primer (100 pmol) (Promega), dNTP-mix (10 mM each) and Transcriptor Reverse Transcriptase (10 units) (Roche). Quantitative real-time PCR was performed in a Bio-Rad iCycler iQ with Bio-Rad iQ SYBR green Supermix. PCR conditions comprised an initial 10 min polymerase activation at 95 °C followed by 35 cycles each consisting of a denaturation step at 95 °C / 15 s and an annealing/elongation step at 60 °C. Used primers were: attacinFW: 5'-TTTTCACAGTCGCACCAATT-3' and attacinREV: 5'-AAACGCCTCTGTCAAATCC-3', defensinFW: 5'-TAGTTTTGGCTTTTCTTACAC-3' and defensinREV: 5'-CGACTACAGTATCCGCTCTTT-3', and cecropinFW: 5'-ATACTCGCTCTTTCAGTCAG-3' and cecropinREV: 5'-CTCTAACAGTAGCGGCAACA-3'; actinFW: 5'-CGCTTCTGGTCTACTACT-3' and actinREV: 5'-CCG-GACATCACAAATGTTGG-3', tubulinFW: 5'-GATGGTCAAGTGGATCCT-3' and tubulinREV: 5'-TGAGAAGTCCCTTCTCC-3'. Three replicates were performed for each sample and threshold cycles (*C<sub>t</sub>*) were recorded and used to calculate gene expression levels.

### 2.6. Statistical data analysis

All statistical analyses were carried out in Stata software version 9.2 ([StataCorp, 2006](#)). A linear regression was used to analyse the loss of weight in individual flies. The difference in weight was used as the response and fly groups (flies emerging from pupae produced by females of the starved and non-starved colony) as discrete explanatory variables. The distribution of the residuals and the heteroskedasticity of the variance were verified ( $p > 0.1$ ).

The proportions of infected flies in different experimental series were compared using a robust logistic regression. Cluster effects resulting from flies infected by feeding on the same mouse and maintained in the same cage were taken into account in the model.

The effect of adult female starvation on the expression levels of attacin, defensin and cecropin in emerging general male flies were analysed using a robust linear model. The response variables were the logarithm of the normalized number of cycles needed for the immune peptides specific cDNA to reach the threshold in real-time PCR. The normalization used was a modification of that proposed by [Vandesompele et al. \(2002\)](#):

$$\text{response} = \ln[(\text{PCR}_{\text{pep}})^{C_{\text{tpep}}}] - \frac{\ln[(\text{PCR}_{\text{act}})^{C_{\text{tact}}}] + \ln[(\text{PCR}_{\text{tub}})^{C_{\text{ttub}}}]}{2}$$

with:

- $\text{PCR}_{\text{pep/act/tub}}$  = PCR efficiency of immune peptides, actin and tubulin, respectively
- $C_{\text{tpep}}$  = number of cycles required for immune peptides genes to reach the threshold in each sample repetition
- $C_{\text{tact/tub}}$  = three repetition average number of cycles required for actin and tubulin genes, respectively to reach the threshold in each sample

Clustering effects resulting from repeated measures of the same samples were taken into account in the model. Discrete explanatory variables consisted of the different experimental groups created in each experiment. For each statistical analysis, differences were considered significant when  $p < 0.05$ .

### 3. Results

During the study period, the starved colony produced 0.07–0.15 pupae per female per week whereas the non-starved colony produced about 1.0 pupae per female per week. Many abortions were observed in the starved colony. The pupal mean weight ranged between 20 and 22 mg in the starved colony while the pupal mean weight in the non-starved colony was about 26 mg. The pupal emergence rate was 89% in the starved colony and 97% in the non-starved female colony.

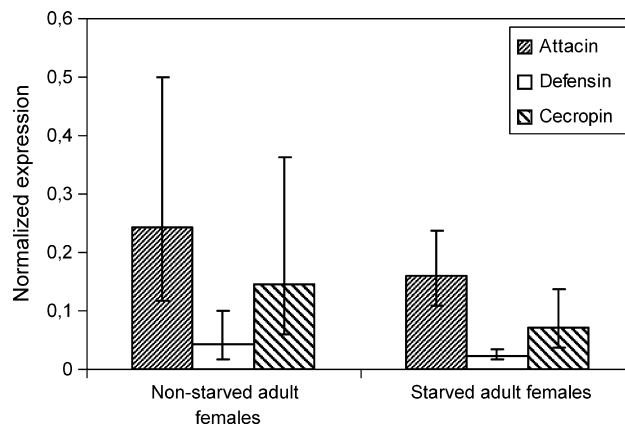
#### 3.1. Nutritional stress and expression of immune peptide genes of the emerging flies

Male flies emerging from pupae produced by the non-starved females had an almost 2-fold higher fat body content than teneral males emerging from pupae produced by the starved females ( $2.5 \pm 0.1$  mg compared to  $1.4 \pm 0.4$  mg,  $p < 0.0001$ ). The normalized gene expression levels of the immune peptides attacin, defensin and cecropin were respectively 1.5, 1.8 and 2.0 times higher in teneral male flies from non-starved adult females (Fig. 1). None of these differences are, however, statistically significant.

#### 3.2. Trypanosome infection rates

The proportion of teneral males feeding on the infected mice was 74.8% and 65.6% for tenerals emerging from pupae produced by the starved and the non-starved females, respectively. Mortality rates of flies after the infective feed were 7.1% and 1.5% for males emerging from pupae produced by starved and non-starved females, respectively.

The proportion of flies from the starved colony that developed a mature infection with *T. congolense* (Table 1) or *T. b. brucei* (Table 2) was significantly higher compared to flies from the non-starved colony ( $p < 0.001$ ). In *T. congolense*-infected flies, the proportion



**Fig. 1.** Normalized expression levels of attacin, defensin and cecropin in male teneral *G. m. morsitans* emerging from non-starved or starved adult female flies and their 95% confidence intervals. Expression level of each immune gene was normalized against the value of the housekeeping genes actin and tubulin.

of male flies that developed a midgut infection was significantly higher in the flies emerging from the starved colony compared to flies from the non-starved colony (0.44 versus 0.27,  $p = 0.0001$ ). The maturation rate of the midgut infections was about 100% in both experimental groups. For infections with *T. b. brucei*, the proportion of flies that developed a midgut infection did not differ significantly between the experimental groups ( $p = 0.067$ ) but the maturation rate was significantly higher in males emerged from pupae produced by females of the starved colony (0.55 versus 0.35;  $p = 0.011$ ) (Table 2).

### 4. Discussion

The starvation induced in the adult female tsetse flies substantially increased their mortality rate and reduced their reproductive

**Table 1**

Proportion of male teneral *G. m. morsitans* emerging from starved or non-starved females and infected with *T. congolense* IL1180.

Experimental group	Batch of flies	Number dissected	Proportion of flies infected in		Maturation rate
			Midgut (immature)	Mouthparts (mature)	
Male teneral flies emerging from starved adult females	1	28	0.50 (14/28)	0.50 (14/28)	1.00
	2	17	0.41 (7/17)	0.41 (7/17)	1.00
	3	25	0.40 (10/25)	0.40 (10/25)	1.00
	4	27	0.41 (11/27)	0.41 (11/27)	1.00
	5	25	0.48 (12/25)	0.44 (11/25)	0.92
	6	17	0.41 (7/17)	0.41 (7/17)	1.00
Male teneral flies emerging from non-starved adult females	1'	15	0.20 (3/15)	0.20 (3/15)	1.00
	2'	22	0.32 (7/22)	0.32 (7/22)	1.00
	3'	40	0.30 (12/40)	0.30 (12/40)	1.00
	4'	46	0.26 (12/46)	0.26 (12/46)	1.00
	5'	28	0.25 (7/28)	0.25 (7/28)	1.00
	6'	31	0.29 (9/31)	0.29 (9/31)	1.00

**Table 2**

Proportion of male teneral *G. m. morsitans* emerging from starved or non-starved females and infected with *T. b. brucei* ANTar 1.

Experimental group	Batch of flies	Number dissected	Proportion of flies infected in		Maturation rate
			Midgut (immature)	Salivary glands (mature)	
Male teneral flies emerging from starved adult females	1	16	0.56 (9/16)	0.25 (4/16)	0.44 (4/9)
	2	20	0.65 (13/20)	0.35 (7/20)	0.54 (7/13)
	3	18	0.61 (11/18)	0.39 (7/18)	0.64 (7/11)
	4	15	0.53 (8/15)	0.33 (5/15)	0.63 (5/8)
	5	28	0.64 (18/28)	0.33 (9/28)	0.50 (9/18)
Male teneral flies emerging from non-starved adult females	1'	20	0.55 (11/20)	0.10 (2/20)	0.18 (2/11)
	2'	26	0.58 (15/26)	0.15 (4/26)	0.27 (4/15)
	3'	25	0.40 (10/25)	0.24 (6/25)	0.60 (6/10)
	4'	27	0.52 (14/27)	0.22 (6/27)	0.43 (6/14)
	5'	43	0.58 (25/43)	0.16 (7/43)	0.28 (7/25)

performance. Moreover, the mortality rate of the offspring of those starved females was substantially higher compared to that of offspring of the non-starved females. This suggests a reduced viability of male tsetse flies produced by female flies submitted to this level of nutritional stress. This is not surprising considering the tsetse's adenotrophic viviparity and the importance of the uterine milk produced by the mother fly for the development of the three larval stages in its uterus (Denlinger and Ma, 1974). Consequently, the condition of the mother fly has direct repercussions for its offspring.

Environmental conditions vary seasonally and have a significant effect on the fat content and size of adult tsetse flies (Glasgow and Bursell, 1960; Van den Bossche and Hargrove, 1999) as well as on the body size and survival of their offspring (Jackson, 1952; Phelps and Clarke, 1974; Dransfield et al., 1989). Based on the performance parameters of the starved colony, the nutritional stress induced in this colony seems to mimic the seasonal stress experienced by field populations of tsetse flies. Especially the reduced fat level of the emerging teneral male flies may be attributed directly to the nutritional stress the female flies were subjected to. In accordance with observations made by Kubi et al. (2006), the mature trypanosomal infection rate of teneral males with low fat levels (emerging from pupae produced by females from the starved colony) was significantly higher than the infection rates of males with higher fat levels (emerging from pupae produced by the non-starved colony). This increased susceptibility to trypanosomal infections could be attributed partially to a reduced baseline immune status of the offspring as a result of the nutritional stress experienced by the parent fly. Such hypothesis is supported by Koella and Sørensen's (2002) observations that sufficient and good quality food is essential for good immune defence in insects. Similarly, Siva-Jothy and Thompson (2002) concluded that a reduction in resource availability could decrease the immune function of invertebrates. In our study, the impact of starvation on the immune system was only examined for immune peptides that are mainly produced systemically by the fat body and used by the tsetse's defence mechanisms against trypanosome infection (Hao et al., 2001; Boulanger et al., 2002). The fact that the expression of these immune peptides genes was reduced, but not significantly so, could be due to the way in which this expression was measured. Indeed, measurement of the expression in the whole abdomen of the fly may mimic significant reductions in the expression of immune peptides in organs where the trypanosomes develop such as the midgut. Other factors that are induced by starvation and that may affect the development of a trypanosomal infection in tsetse can, however, not be excluded. They include midgut lectins, anti-oxidants and symbiotic associations in tsetse (Abubakar et al., 2006; Munks et al., 2005; Macleod et al., 2007; Roditi and Lehane, 2008).

Although these observations are based entirely on laboratory experiments, the outcome may contribute to a better understanding of the seasonal dynamics of trypanosomal infection rates in field populations of tsetse flies. Indeed, the monthly metacyclic infection rate of tsetse populations does undergo substantial variations over the year. Such variations have been attributed to changes in the survival rate of the tsetse population or the availability of infected hosts (Jordan, 1965; Harley, 1967). Our results suggest that seasonal stress situations, resulting in changes in the susceptibility to trypanosomal infections of emerging teneral flies, could also play a major role. Especially the hot dry season when ambient temperatures are high constitutes a stressful period for tsetse flies. This is reflected in the decrease in the fly's body size and fat body level during this season (Jackson, 1952; Glasgow and Bursell, 1960; Dransfield et al., 1989). The association between high ambient temperatures, stress in tsetse flies and the concomitant increase in the tsetse's susceptibility to trypanosomal infections may offer an explanation for the often observed relationship between high temperatures and high trypanosomal infection rates in tsetse. For example, Kinghorn

and York, 1912 and Burt (1946) reported higher susceptibility to infection with *T. b. rhodesiense* of flies emerging from pupae incubated at a high temperature. The same association may explain partly Fairbairn's (1948) observation that in the western *G. morsitans* belt of Tanzania the monthly mean maximum temperature and the number of diagnosed cases of sleeping sickness was significantly correlated. Finally and on a continental scale, a review of surveys determining the infection rates of savannah species of tsetse flies conducted over a period of 45 years also showed a positive correlation between infection rate of tsetse flies and the mean annual temperature (Ford and Leggate, 1961). Nevertheless, further research is required to determine the field implications of our laboratory observations.

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