Antiviral drug valacyclovir treatment combined with a clean feeding system enhances the suppression of salivary gland hypertrophy in laboratory colonies of Glossina pallidipes

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Abstract

**Background:** Hytrosaviridae cause salivary gland hypertrophy (SGH) syndrome in some infected tsetse flies (Diptera: Glossinidae). Infected male and female G. pallidipes with SGH have a reduced fecundity and fertility. Due to the deleterious impact of the virus on G. pallidipes colonies, adding the antiviral drug valacyclovir to the blood diet and changing the feeding regime to a clean feeding system (each fly receives for each feeding a fresh clean blood meal) have been investigated to develop virus management strategies. Although both approaches used alone successfully reduced the virus load and the SGH prevalence in small experimental groups, considerable time was needed to obtain the desired SGH reduction and both systems were only demonstrated with colonies that had a low initial virus prevalence (SGH ≤ 10%). As problems with SGH are often only recognized once the incidence is already high, it was necessary to demonstrate that this combination would also work for high prevalence colonies.

**Findings:** Combining both methods at colony level successfully suppressed the SGH in G. pallidipes colonies that had a high initial virus prevalence (average SGH of 24%). Six months after starting the combined treatment SGH symptoms were eliminated from the treated colony, in contrast to 28 months required to obtain the same results using clean feeding alone and 21 months using antiviral drug alone.

**Conclusions:** Combining valacyclovir treatment with the clean feeding system provides faster control of SGH in tsetse than either method alone and is effective even when the initial SGH prevalence is high.

**Keywords:** Diptera, Glossinidae, Salivary gland hypertrophy virus, Hytrosaviridae

Findings

Tsetse flies (Glossina spp.) are the vectors of sleeping sickness in humans (human African trypanosomosis), and the cattle disease nagana [1,2]. Nagana is considered among the greatest constraints to livestock production in sub-Saharan Africa. The prophylactic and curative treatment of livestock with trypanocidal drugs is problematic due to the development of resistance of the parasites to the available trypanocidal drugs. Therefore, removal of the vector, the tsetse fly, is considered the most efficient way of managing nagana [3,4].

There are a number of efficient tsetse control tactics available that can be combined and applied in the frame of the area-wide integrated pest management approach. One of these tactics is the sterile insect technique (SIT), which is very efficient for controlling and eliminating low-density populations previously reduced with other control methods [5]. Using this approach, a successful sustainable eradication of Glossina austeni from Unguja Island (Zanzibar), United Republic of Tanzania, was achieved [6]. Consequently, other programmes were developed to apply this approach on the African mainland and, in 1996, the Government of Ethiopia embarked on such a programme to create a zone free of Glossina pallidipes in the Southern Rift Valley of Ethiopia [7,8]. In this programme the establishment and expansion of

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G. pallidipes colonies for mass-production of sterile male flies proved to be difficult and several colony collapses were experienced. The cause of these failures was identified as the salivary gland hypertrophy virus (GpSGHV), which was often present at high prevalence and affects colony productivity and stability [9].

The virus transmission mode in wild tsetse populations is most probably mother to offspring transmission, either trans-ovum or through infected milk glands. In laboratory-maintained flies where blood feeding using an in vitro membrane feeding system is used, horizontal transmission was found to be the most important route of virus transmission. In order to develop GpSGHV management strategies several studies were conducted on various aspects of the biology of the virus [10-13]. Following these basic studies, several virus management approaches were suggested [14] and the effect of two of these strategies on the prevalence of SGH in colonies of G. pallidipes was recently published i.e. adding the antiviral drug valacyclovir to the blood diet of the flies [15,16] and the application of a clean feeding strategy (each fly received a clean fresh blood meal each time) [17,18]. Briefly, the clean feeding system can be initiated in a large-scale colony by selecting as many cages of teneral flies from the main colony as can be fed at one time on the available clean membranes and thereafter these flies and their progeny should always be fed first on the fresh, clean blood (that has not been supplied to other flies). These flies will, over time develop into a clean feeding colony, and when the virus reaches a very low level (no more hypertrophy) so that contamination of the blood is minimal, a second set of cages of teneral flies is started in the same manner, fed second on the membranes now containing essentially no virus. This process is continued group by group until all the flies are kept in these separate groups. Both clean feeding and the addition of valacyclovir successfully reduced the virus load and the prevalence of SGH symptoms in the treated G. pallidipes groups when implemented separately. More rapid results were obtained when these approaches were combined with selection of low virus prevalence at a small scale experimental level [18]. Despite the successful reduction of the SGH prevalence in these colonies, both approaches had some obvious limitations i.e. (i) the long time needed after treatment initiation to achieve significant reduction in virus load (21 months with valacyclovir and 28 months with clean feeding system) and (ii) both approaches were only demonstrated for groups with a low prevalence of SGH symptoms.
initial virus load and low SGH prevalence (≤10%). In operational large scale production facilities, the detection of virus problems usually surfaces when a lower productivity of the colony becomes apparent, usually when the prevalence of the virus is already high. In such cases, the main objective is to restore the productivity of the colony (i.e. reduce the virus load) in the shortest possible time to prevent any interruption of the operational program. We therefore tested the impact of the combined approaches in large scale colony level on a high virus prevalence colony.

All experiments were carried out with the colony described previously by Abd-Alla et al. [19], kept in the same manner. Previous analysis revealed that ca 4–10% of individuals in this colony showed SGH [18]. A high SGH prevalence colony (SGH of 24%) was developed by exposing flies to the virus by feeding blood contaminated with the virus. Flies from this colony were used to start two new experimental colonies: the first to be maintained under a clean feeding system combined with valacyclovir treatment (the flies were fed on fresh blood supplemented with valacyclovir (300 μg/ml)) [18], the flies of the second colony were fed on contaminated blood. The blood for this colony was first used for two feeding rounds of flies of the main colony, then the blood was collected from the feeding trays and valacyclovir was added to the required concentration (300 μg/ml) and used for feeding the experimental flies. The prevalence of SGH was determined by dissection of flies (n = 360) from different ages.

The graphics were created using the ggplot2 package [20] in R [21]. Dissection results were analysed by analysis of variance (ANOVA/ANCOVA) in R [20] after arc-sine square root transformation.

The dissection results presented in Figure 1 show that the prevalence of the SGH syndrome in the clean feeding system with valacyclovir treatment was significantly reduced ($F_{2,64} = 93.989, P < 0.001$) after six months i.e. from 24% (the average prevalence in the colony before treatment [21]) to 0%, irrespective of the age of the flies ($F_{1,64} = 0.914, P = 0.343$). In the valacyclovir treatment with contaminated feeding, the prevalence of SGH (1.4%) was significantly reduced after six months ($F_{1,24} = 73.304, P < 0.001$) but the SGH was only eliminated from the treated flies after 21 months. Although the difference in the prevalence of SGH between the valacyclovir treatment implemented alone or combined with clean feeding at 6 or 21 months was not significant ($F_{1,52} = 2.634, P = 0.111$), the antiviral drug without the clean feeding system required 21 months post treatment to achieve complete elimination of SGH syndrome from the treated colony.

Taken together, the data presented in this paper demonstrate that the clean feeding strategy to reduce the virus load and to remove the SGH syndrome in colonies of G. pallidipes can be improved by combining this system with a valacyclovir treatment. The time needed to eliminate the SGH syndrome from the colony was reduced from 28 months [19] to 6 months, leading to sustainable maintenance and expansion of the colony. The data together with Abd-Alla et al. [19] also confirm that the implementation of the feeding regime in combination with valacyclovir treatment is readily applicable in tsetse mass rearing facilities and only requires changes to the colony handling and recording system, which only needs minimal additional training of the staff at no significant additional cost to the SIT program compared to the cost of other methods to manage the virus. Also, once the prevalence of SGH has been reduced to zero, antiviral drug treatment may be withdrawn to reduce the cost so long as the clean feeding system is strictly observed. The data presented in this article strongly supports the implementation of a clean feeding strategy in combination with valacyclovir treatment in large-scale G. pallidipes production facilities in order to achieve sustainable GpSGHV control.

**Abbreviations**

GpSGHV: Glossina pallidipes salivary gland hypertrophy virus; SGH: Salivary gland hypertrophy; SIT: Sterile insect technique.

**Competing interests**

The authors declare that they have no competing interests.

**Authors’ contributions**

Conceived and designed the experiments: AMMA, AGP and MJBV. Performed the experiments: CM. Drafted the manuscript: AMMA, AGP and MJBV. All authors read and approved the final manuscript.

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