

MOLECULAR ANALYSIS OF GUT CONTENTS TO ESTABLISH HOST RANGE OF EDIBLE GRASSHOPPERS IN EAST AFRICA

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INTRODUCTION

- African grasshopper (*Acanthacris ruficornis*) and long-horned grasshoppers (*Ruspolia differens*) are important cultural food among communities in Kenya and Uganda¹.
- These edible insects are a good source of protein, fats, vital minerals and vitamins.
- Unfortunately, communities rely on wild grasshoppers that only multiply during the rainy season.
- Despite their traditional importance as food, mass-rearing protocols for the species has been undeveloped².
- Need to develop mass rearing protocols for edible grasshoppers to enhance food security in East Africa exists.
- To optimise the mass-rearing protocol, information on the preferred host plants of the grasshopper is critical.

OBJECTIVE

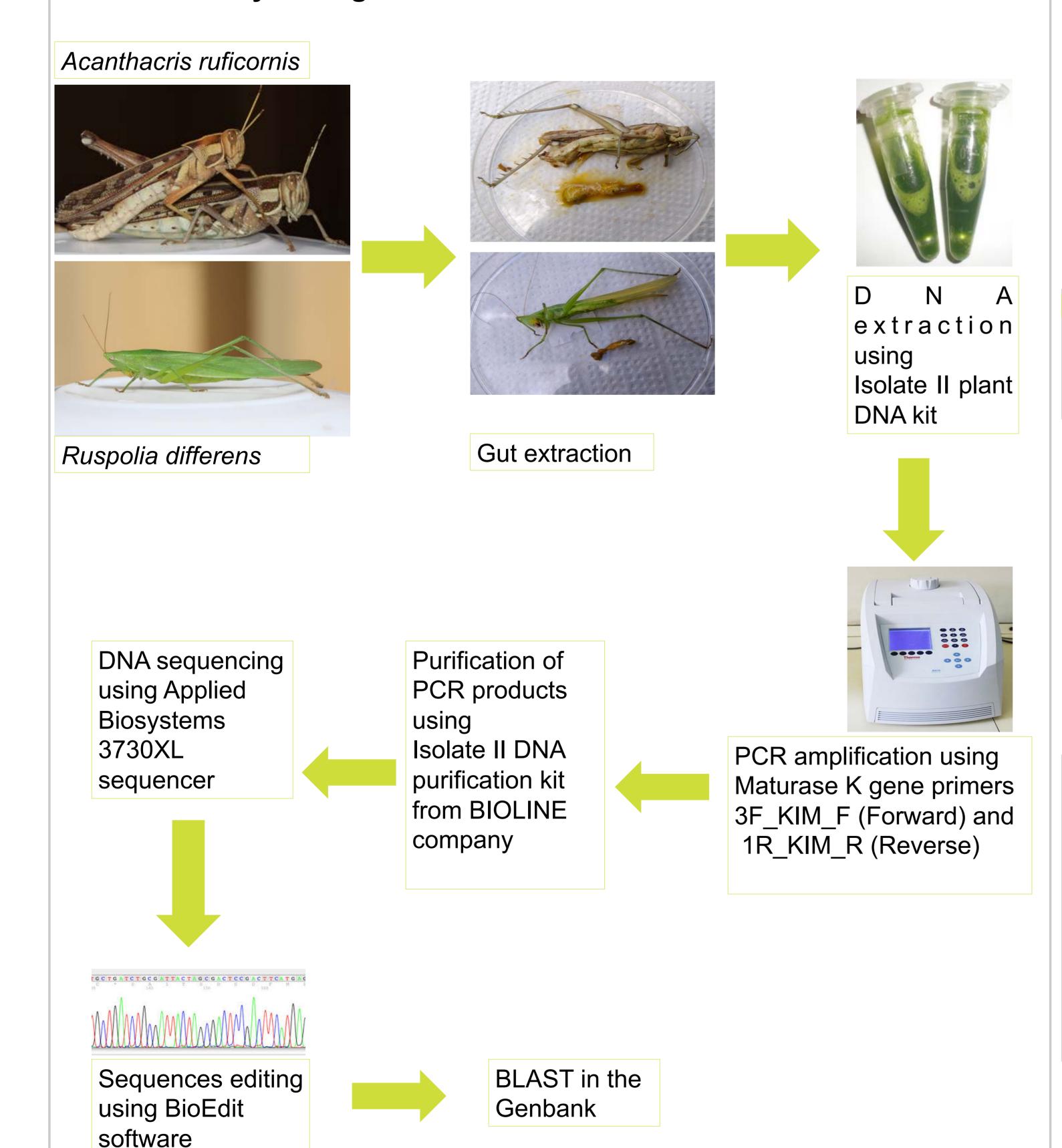
To determine host plants of *Ruspolia differens* in Uganda and *Acanthacris* ruficornis in Kenya through molecular analysis of gut contents.

METHODS

Sampling

Ruspolia differens were collected using light traps in Mbarara, Kabale and Hoima, Uganda. Acanthacris ruficornis were collected in Murang'a, Kenya using sweep nets. Samples were preserved in absolute ethanol until further analysis.

Molecular analysis of gut content³





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RESULTS

Table 1: Outcome of the gut content analysis to identify host plants of Ruspolia differens and Acanthacris ruficornis

Site	Grasshopper	Host family	Host species	GenBank Accessions with Similarity (%)
Murang'a	Acanthacris ruficornis	Amaranthaceae	Achyropsis Ieptostachya	AY998117.1 (99)
		Tiliaceae	Heliocarpus pallidus	KM219813.1 (97)
		Apiaceae	Centella virgata	KP110015.1 (99)
		Asteraceae	Erigeron atticus	KP175136.1 (87)
		Poaceae	Digitaria exilis	KJ513091.1 (99)
		Nyctaginaceae	Bougainvillea glabra	JQ844141.1 (100)
Mbarara	Ruspolia differens	Poaceae	Digitaria exilis	KJ513091.1 (99)
Hoima	Ruspolia differens	Poaceae	Digitaria exilis	KJ513091.1 (99)
Kabale	Ruspolia differens	Poaceae	Digitaria exilis	KJ513091.1 (99)

CONCLUSION

- Molecular analysis of gut contents has identified the host range of the two edible grasshoppers.
- Digitaria exilis was the only identified host plant of Ruspolia differens.
- Six plants were identified as hosts of *Acanthacris ruficornis*, all of which were broad-leaved species except *D. exilis* which is a grass species.
- The identified wild host plants of these insects are available in most parts of East Africa; hence, they can be adopted for mass rearing of the two edible grasshoppers.

IMPACT

The identified host plants of *Ruspolia differens* and *Acanthacris ruficornis* will lead to more efficient and sustainable mass-rearing of these grasshoppers.

REFERENCES

- 1. Agea, G., Biryomumaisho, D., Buyinza, M. & Nabanoga, G. (2008). *African Journal of Food, Agriculture, Nutrition and Development*, 8, 291–303.
- 2. Malinga, G. M., Valtonen, A., Lehtovaara, V. J., Rutaro, K., Opoke, R., Nyeko, P. & Roininen, H. (2018). *Applied Entomology and Zoology*, 53, 229–236.
- 3. Avanesyan, A. (2014). Applications in Plant Sciences, 2(2), 1300082.

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