Proceedings of the XIII International Symposium on Biological Control of Weeds

September 11–16, 2011 Waikoloa, Hawaii, USA

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CONTENTS

PREFACE	xxv
INTRODUCTION	
Symposium Welcome T. Johnson and P. Conant	xxix
Opening Address: The future challenges of invasive species work <i>W. W. M. Steiner</i>	xxx
SESSION 1: PRE-RELEASE TESTING OF WEED BIOLOGICAL CONTROL AGENTS	
Papers	
Pre-release studies and release of the grasshopper <i>Cornops aquaticum</i> in South Africa – a new biological control agent for water hyacinth, <i>Eichhornia crassipes</i> A. Bownes, A. King and A. Nongogo	3
Australia's newest quarantine for weed biological control W. A. Palmer, T. A. Heard, B. Duffield and K. A. D. W. Senaratne	14
Host specificity of an Italian population of <i>Cosmobaris scolopacea</i> (Coleoptera: Curculionidae), candidate for the biological control of <i>Salsola tragus</i> (Chenopodiaceae) M. Cristofaro, F. Lecce, A. Paolini, F. Di Cristina, MC. Bon, E. Colonnelli and L. Smith	20
Biological control of Chilean needle grass (<i>Nassella neesiana</i> , Poaceae) in Australasia: Completion of host range testing <i>F. Anderson, L. Gallego, J. Barton and D. McLaren</i>	26
Abstracts	
Finding the weapons of biomass destruction — identifying potential biological control agents by applying principles of chemical co-evolution <i>M. R. Berenbaum</i>	33
Molecular analysis of host-specificity in plant-feeding insects: Phylogenetics and phylogeography of Fergusonina flies on Australian paperbarks S. Scheffer, R. Giblin-Davis, M. Purcell, K. Davies, G. Taylor and T. D. Center	of 34
Selection of test plant lists for weed biological control with molecular and biochemical data <i>G. S. Wheeler</i>	35
Successfully eliminating parasitic gregarines from <i>Neolema ogloblini</i> (Coleoptera: Chrysomelidae) — a biological control agent for <i>Tradescantia fluminensis</i> (Commelinaceae) L. A. Smith, S. V. Fowler, Q. Paynter, J. H. Pedrosa-Macedo and P. Wigley	36

Effects of drought on the biological control of spotted knapweed Y. K. Ortega and D. E. Pearson	309
Solanum elaeagnifolium (Solanaceae), an alien invasive weed for Greece and southern Europe, and newly discovered endemic natural enemies	
J. Kashefi, G. Ara, W. Jones and D. Strickman	310
Microsatellites uncover multiple introductions of clonal giant reed (<i>Arundo donax</i>) in the new wo D. Tarin, A. E. Pepper, J. Goolsby, P. Moran, A. C. Arquieta, A. Kirk and J. R. Manhart	orld 311
Utility of microsatellite markers from the wheat genetic map in the genome of medusahead rye (<i>Taeniatherum caput-medusae</i>) B. G. Rector, M. C. Ashley and W. S. Longland	312
The interaction between drought and herbivory by a biological control agent on populations of th invasive shrub <i>Tamarix</i> sp.	e
W. I. Williams and A. P. Norton	313
Doct introduction evalution in the higherinal control execut I question we incohere	
Post-introduction evolution in the biological control agent <i>Longitarsus jacobaeae</i> M. Szűcs, U. Schaffner and M. Schwarzländer	314
Eurasian watermilfoil phenology and endophyte abundance and diversity J. F. Shearer, M. J. Grodowitz and B. D. Durham	315
Herbivore-induced plant defenses and biological control of invasive plants <i>J. B. Runyon and J. L. Birdsall</i>	316
Comparison of native and invasive populations of <i>Taeniatherum caput-medusae</i> ssp. <i>asperum</i> (medusahead): Evidence for multiple introductions, source populations and founder effects	
M. Peters, R. Sforza and S. J. Novak	317
Morphological and genetic differentiation among subspecies of <i>Taeniatherum caput-medusae</i> : Disentangling taxonomic complexity in the native range	
M. Peters, R. Sforza and S. J. Novak	318
District souther of Australia Australia and successful a Laurence for the same	
Biological control of <i>Ambrosia artemisiifolia</i> : Learning from the past <i>H. Müller-Schärer and U. Schaffner</i>	319
Effect of nitrogen addition on population establishment of the Arundo armored scale Rhizaspidio	itus
donacis	220
P. J. Moran and J. A. Goolsby	320
Stenopelmus rufinasus proves to be an excellent Azolla taxonomist	
M. Hill and P. Madeira	321
What do chloroplast sequences tell us about the identity of Guinea grass, an invasive Poaceae in the	he
southern United States?	225
MC. Bon, J. Goolsby, G. Mercadier, T. Le Bourgeois, P. Poilecot, M. Jeanneau and A. Kirk	322



What do Chloroplast Sequences Tell us about the Identity of Guinea Grass, an Invasive Poaceae in the Southern United States?

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Abstract

The commonly named Guinea grass of the Poaceace family is a native African grass that has been extensively and successfully introduced as a source of animal fodder to other tropical areas of both hemispheres. On a global scale but particularly in the southern United States, the Caribbean and Hawaii, it is becoming a serious threat to biodiversity not only due to its invasiveness but also because it produces high fuel loads for fires. For the first time, a biological control program is being attempted in Texas. Source populations of the Texan invasion have to be identified in the native range in order to facilitate the search for potential biological control agents. This raises the critical issue of a proper taxonomic identification for this taxon with a history of taxonomic revisions, multiple scenarios of massive introductions and hybridization and polyploidisation events. Guinea grass in the strict sense should refer to Megathyrsus maximus (Jacq.), also known as Panicum maximum and Urochloa maxima. To unravel the taxonomic identification and the evolutionary history of this controversial taxon, we have begun to analyze sequences of two chloroplast regions in modern African and Texan samples as well as historical specimens in the CIRAD collection, some dating back to 1944, prior all extensive improvement programs in Africa. None of the sequences matched the sequence of a voucher specimen of Megathyrsus maximus (Jacq.). Results provided evidence of two different maternal lineages, one distributed from eastern Africa to southeastern Africa and Texas that fully matched the sequence of a voucher specimen of Megathyrsus infestus (Andersson) and one distributed across western/central Africa and French Guiana that do not belong to Megathyrsus genus. Future programs of exploration and collection of natural enemies are to be reviewed in light of these findings.