Response of Cucurbit Species to *Rotylenchulus reniformis*

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ABSTRACT

The state of Rio Grande do Norte is the major melon (*Cucumis melo*) producer in Brazil and the reniform nematode *Rotylenchulus reniformis* has become increasingly important due to damages to that crop and the lack of resistant cultivars and effective control measures. Under these circumstances, eight cucurbit genotypes were screened for resistance or tolerance to a population of *R. reniformis* in naturally infested field soil, under greenhouse conditions. A high reduction in shoot mass was found in all infected genotypes. Watermelon (*Citrullus lanatus*) cv. Sugar Baby showed the lowest rate of nematode reproduction.

Additional keywords: reniform nematode, resistance, tolerance, *Cucumis* spp.

RESUMO

Resposta de genótipos de cucurbitáceas a *Rotylenchulus reniformis*

O estado do Rio Grande do Norte é o principal produtor de melão (*Cucumis melo*) do Brasil, tendo o nematóide *Rotylenchulus reniformis* relevante importância devido ao potencial de dano à cultura e inexistência de cultivares resistentes e medidas de controle efetivas. Sob estas circunstâncias, oito espécies de cucurbitáceas foram avaliadas quanto a resistência e tolerância a uma população nativa de *R. reniformis* em solo naturalmente infestado, em casa de vegetação. Ocorreu alta redução em biomassa fresca da parte aérea em todos os genótipos estudados. A melancia (*Citrullus lanatus*) cv. Sugar Baby apresentou a mais baixa taxa de reprodução do nematóide.


From 1996 to 2001, the state of Rio Grande do Norte yielded 57% of the Brazilian melon (*Cucumis melo L.*) production (IBGE, 2003). In Mossoró and Açu municipalities, high yield losses have recently been associated with high population densities of the reniform nematode (*Rotylenchulus reniformis* Linford & Oliveira) (Moura et al., 2002). A lack of resistant genotypes for practical and effective nematode control measures has limited profitable melon yields in these areas. The objective of this research was to evaluate, in the greenhouse using naturally infested soil, the reaction of cucurbit genotypes (seven local and economically valuable crops) along with an ordinary weed) to the reniform nematode. This information is needed for management assistance in crop rotation. The genotypes evaluated were the melons *C. melo* var. *cantalupensis* Naud. and *C. melo* var. *inodorus* Naud., cucumber (*Cucumis sativus* L.) cv. Marketer, gherkin (*Cucumis anguria* L.) cv. Maxixe do Norte, watermelon (*Citrullus lanatus* Thunb.) cv. Sugar Baby, summer squash (*Cucurbita pepo* L.) var. *melopepo* cv. Abobrinha Brasileira, squash (*Cucurbita moschata* Duch. ex Poir.) cv. Baiana Tropical and the weed *Mordomica charantia* L. The experiment was composed of 16 treatments (eight genotypes in both naturally infested soil and control) and four replications arranged in a completely randomized design. Each replication corresponded to a plant grown in a 500-cm³ pot, filled with naturally infested soil from Baraúna municipality melon fields. Nematode initial population (*Pi*) was assessed by centrifugal flotation procedure according to Jenkins (1964), and determined to be around five juveniles and immature females forms/500 cm³ of soil. Evaluations were carried out 70 days after seeds were sown, based on the relative biomass (RB) of each germplasm. The RB was the shoot fresh mass (SFM) yielded in the presence of the nematode divided by SFM in the absence of it. Reproduction factor (RF = *Pi*/*Pi*) was calculated using infective nematodes in the root system plus juveniles, immature females and males in soil as the final population (PF). Infective nematode stages were recorded staining root systems with acid fuchsin (Byrd et al., 1983). Data were submitted to analysis of variance using the SAS GLM procedure and mean separation with Tukey’s multiple range test (*P = 0.05*). Reproduction data (x) were transformed to log10(x+1) values prior statistical analysis and reported as antilogs.

Although RB values ranged from 0.32 (*C. melo* var. *inodorus*) to 0.64 (*C. pepo* var. *melopepo*), there was no difference (*P > 0.05*) among genotypes (Figure 1A). In contrast,
RF was affected ($P=0.05$) by genotype (Figure 1B). Reproductive factors ranged from 1 to 98, with *C. lanatus* cv. Sugar Baby, tending to suppress the nematode reproduction. Due to the high reproduction of *R. reniformis* in *C. melo* var. *inodorus*, this genotype is not recommended for infested areas. This fact suggests that after a few cropping seasons the density of nematode may increase to damage levels, therefore inducing high economic losses. This is also true for *M. charantia*, *C. moschata* cv. Baiana Tropical, *C. pepo* var. *melopepo* cv. Abobrinha Brasileira, *C. anguria* cv. Maxixe do Norte and *C. melo* var. *cantalupensis* (Figure 1B). *Momordica charantia*, a common weed in melon fields, presented high RF, indicating that this species must be eradicated in melon areas.

In Nematology, resistance has been defined as the effect of the host on nematode reproduction, and tolerance as the absence of yield suppression due to the nematode parasitism (Cook & Evans, 1987). The search for resistance to *R. reniformis* has been carried out mostly on cotton (*Gossypium hirsutum* L.) and soybean (*Glycine max* L.) in
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At the moment, there are no reniform nematode-resistant (or tolerant) melon genotypes available for use in Brazil. However, a watermelon genotype has been found to suppress nematode reproduction, although plant development has also been suppressed. The final population level is critical for management strategies because it may affect crop-rotation sequences. As watermelon is one of the major sources of agricultural income in that state, its introduction in rotation with melon could be useful but is often limited by the lack of alternative crops of equality high-value. Field studies are recommended for more accurate performance of the watermelon Sugar Baby genotype, including yield suppression in highly infested soil.

**LITERATURE CITED**


