

# Rice Root-Knot Nematode *Meloidogyne graminicola* (Nematoda: Chromadorea: Tylenchida: Meloidogynidae: Meloidogyne)<sup>1</sup>

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## Introduction

Rice root-knot disease is caused by different *Meloidogyne* species (*Meloidogyne graminicola*, *M. hainanensis*, *M. incognita*, *M. javanica*, *M. arenaria*, *M. oryzae*, *M. salasi* and *M. tryticoryzae*). However, *Meloidogyne graminicola* is considered the most damaging root-knot species to Asian rice cultivation due to its ability to survive under flooded soil conditions (Bridge et al., 2005; Mantelin et al., 2017). This nematode has been found in rice nurseries, rainfed upland rice and lowland rice but is also widespread in deep-water, irrigated rice production systems. Yield loss due to *Meloidogyne graminicola* ranges from 28%–87% depending on disease severity and cultivar (Bellafiore et al., 2015).

## Distribution

*Meloidogyne graminicola* has been reported in upland, irrigated, lowland and deep-water rice in South and Southeast Asia. It is also found in the United States and Latin America, and was recently reported in Africa and Europe (Bridge et al., 2005; Dutta et al., 2012; Kyndt et al., 2014; Mantelin et al., 2017; Sacchi et al., 2021).

In the United States, *Meloidogyne graminicola* was first reported in Stuttgart, Arkansas in 1934 and then in Baton Rouge, Louisiana in 1965 (Yik and Birchfield, 1979). Since then, *Meloidogyne graminicola* has been reported to occur in other states like Texas, Georgia, and Mississippi (MacGowan and Langdon, 1989; EPPO, 2017).

## Symptoms and Plant Part Affected

Yellowing, stunting, and hook-like galls on the roots of rice plants are the characteristic symptoms caused by *Meloidogyne graminicola* (Figure 1). Severely infected plants flower and mature earlier than healthy plants. Distortion and crinkling along the margins of the newly emerged leaves may also indicate infection (Bridge et al., 2005; Kyndt et al., 2014).

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Figure 1. Typical hook-like galls on rice roots infected by *Meloidogyne graminicola*.

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## Lifecycle (Figure 2)

*Meloidogyne graminicola*, like other *Meloidogyne* species, develops from an embryo through the first and second juvenile stages within an eggshell (Figure 3). The infective second-stage juvenile (J2s) hatches and (Figure 4) infects at the elongation zone and then moves upward to the root tips where they invade the vascular cylinder to form a feeding site of three to ten cells, called a giant cell. Simultaneously, the neighboring cells start to divide to form a typical gall or root-knot. Male and female J3s become round and sedentary inside the gall and continuously molt to J4s and adults (Figure 5). *Meloidogyne graminicola* females commonly reproduce asexually through parthenogenesis. What makes *Meloidogyne graminicola* different from other *Meloidogyne* species is that females lay eggs typically within host tissues as an adaptation to the flooded conditions found in rice fields. The newly hatched J2s (Figure 4) can stay inside the gall or move intercellularly to establish a new giant cell within the same root. In rare cases (c.a 0.5%), *Meloidogyne graminicola* reproduces through sexual reproduction in which adult males remodel back to a vermiform shape, while adult females keep their pear-shape. Adult males fertilize females and leave the root afterward. The life cycle of *Meloidogyne graminicola* can be completed in as few as 19 days under ideal environmental conditions. A single female can lay 500 to 1000 eggs during its lifetime (Bridge et al., 2005; Kyndt et al., 2014; Mantelin et al., 2017).

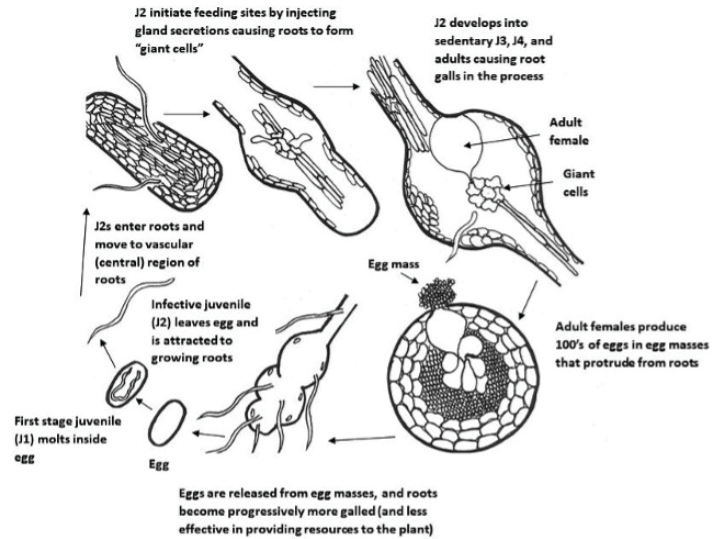


Figure 2. Generalized life cycle of the root-knot nematode *Meloidogyne* spp. (courtesy H. Regier, adapted from G. Abawi and V. Brewster). Note that, unlike other *Meloidogyne* species, *Meloidogyne graminicola* lays eggs within the host as an adaptation to flooded conditions.



Figure 3. Root-knot nematode first-stage juvenile (J1) in the eggshell. Note that this image is of *Meloidogyne incognita*.

Credits: Hung Xuan Bui, UF/IFAS

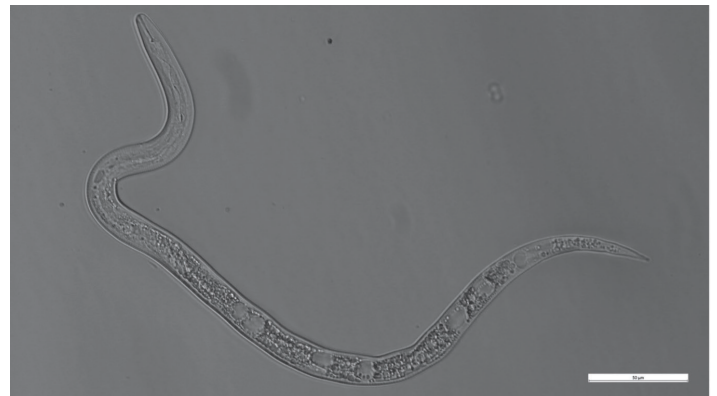


Figure 4. Root-knot nematode second-stage juvenile (J2) initially moves in the soil to find the root to invade in the early season. Note that this image is of *Meloidogyne javanica*.

Credits: Hung Xuan Bui, UF/IFAS



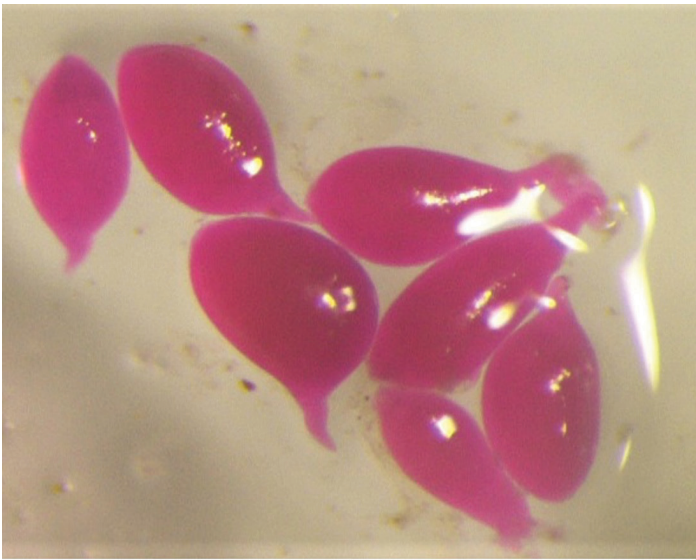


Figure 5. Following infection and the onset of feeding, *Meloidogyne graminicola* develops from a vermiform J2 into a pear-shaped adult. These individuals were stained with a red dye.

Credits: Hung Xuan Bui, UF/IFAS

## Diagnosis

*Meloidogyne graminicola* induces the formation of characteristic hooked-like galls, normally at the root tips (Figure 1 & 7). Disease severity can be evaluated based on a root-knot galling index for rice root-knot (Figure 8). In the field, symptoms can be observed as heavily reduced growth, empty grains, reduced tillering, chlorosis, and wilting, all of which will typically occur in patches (Figure 7) (Bridge et al., 2005; Kyndt et al., 2014; Mantelin et al., 2017).

To confirm the infection on rice is due to *Meloidogyne graminicola*, soil and root samples should be sent to a nematology laboratory for identification. Perineal patterns of adult females are often used as a diagnostic procedure; however, this approach can lead to misidentification due to the overlapping morphological characters with other *Meloidogyne* species (Chen et al., 2019; Trinh et al., 2019). Molecular identification now is more rapid and accurate for root-knot nematode species identification by both species-specific primers and sequence information of conserved regions (Htay et al., 2016; Chen et al., 2019).

For instance, our nematology laboratory molecularly identified *Meloidogyne graminicola* collected from a field heavily infested with nutsedge in 2019 at the UF/IFAS Gulf Coast Research and Education Center (UF/IFAS GCREC), Wimauma, Florida. The identification was done by using D2A/D3B primer set. DNA extraction, polymerase chain reaction, DNA sequencing, and phylogenetic analysis as described by Oliveira et al. (2019) (Figure 6). *Meloidogyne graminicola* is identified in sandy soil located in GCREC.

The DNA sequences were deposited in the GeneBank database under accession numbers: MZ151167 and MZ151168.

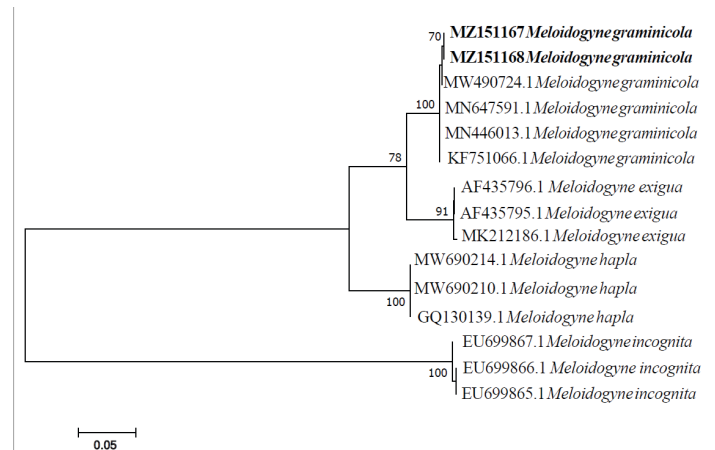


Figure 6. Phylogenetic analysis within populations and species of the genus *Meloidogyne* as inferred from maximum likelihood analysis using the D2-D3 of 28S rRNA gene sequence dataset with T92+G model. Newly obtained sequences are indicated in bold and the others were obtained from the GeneBank database.

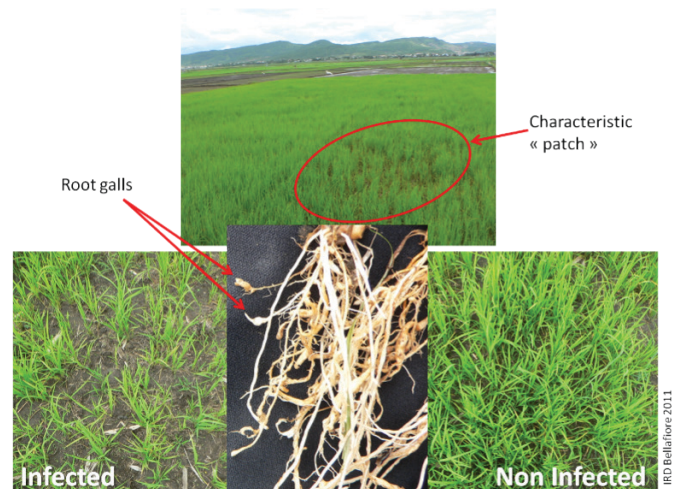


Figure 7. Rice field infected by *Meloidogyne graminicola* with the typical patches of reduced growth and tillering symptoms and roots with hooked-like galls.

Credits: Bellaïre, PHIM Plant Health Institute, University Montpellier, IRD, CIRAD, INRAE, Institute Agro, Montpellier, France

## Hosts

The main host is rice, but *Meloidogyne graminicola* has a very wide host range including wheat, barley, sorghum, soybean, okra, green gram, berseem, potato, onion, garlic and weed species like *Cyperus rotundus*, *Echinochloa colonum*, *E. crusgalli*, *Leptochloa colonicus* and *Phalaris minor* (Bridge et al., 2005; Mantelin et al., 2017). In Florida, the nematode was found on purple nutsedge in Wimauma, FL in 2019 and on sandbur weed (*Cenchrus* spp.) (Handoo et al., 2003).

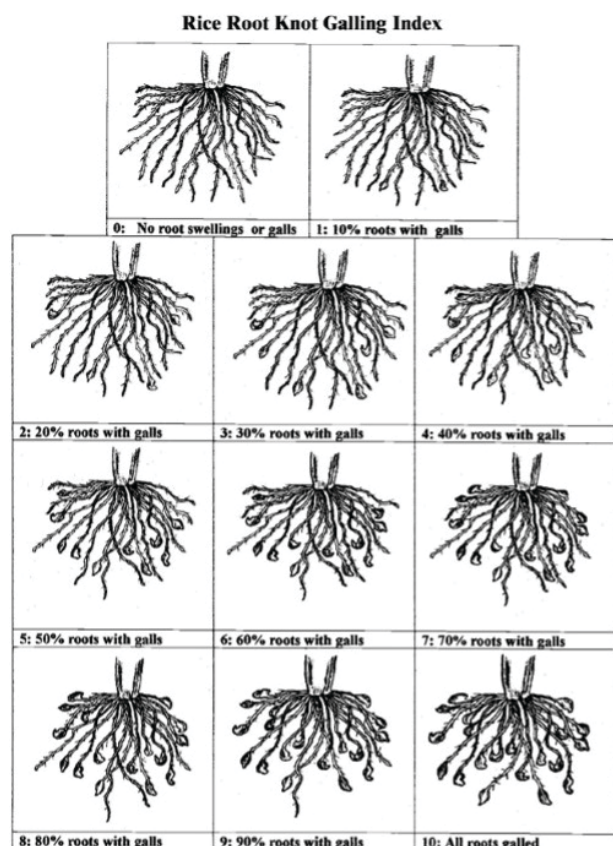


Figure 8. Rice root-knot nematode gall index for evaluating the infection severity caused by *Meloidogyne graminicola*. Credits: Adapted from Bridge et al. (2005)

## History and Economic Importance

In 1934, rice root-knot nematode was first reported to parasitize rice (*Oryzae sativa*) in Arkansas and was named *Heterodera marioni*, which was later renamed to *Meloidogyne graminicola* by Golden and Birchfield (Yik and Birchfield, 1979). *Meloidogyne graminicola* can cause yield loss in rice of 10 to 80% under heavy infestations. Also, its ability to adapt to deep-water conditions makes this nematode a threat to all types of rice agrosystems (Kyndt et al., 2014; Mantellin et al., 2017).

In Florida, rice is cultivated mainly in the Everglades Agricultural Area with a steady increase from 11,912 acres in 2008 to approximately 29,000 acres in 2017 (Bhadha et al., 2018). To our knowledge there have been no reports of rice root-knot nematode from the Everglades Agricultural Area.

## Management

### Cultural Strategies

Although *Meloidogyne graminicola* can survive and reproduce under flooded conditions, continuous flooding from seedling to maturity can help prevent the infection of

new rice roots. Crop rotation with non-host plants such as sweet potato, cowpea, sesame, castor, sunflower, soybean, turnip, cauliflower, jute, mustard and chickpea for at least 12 months are recommended to help manage rice root-knot nematode (Dutta et al., 2012; Mantellin et al., 2017).

### Host Resistance

*Meloidogyne graminicola* can infect most of the commercial rice cultivars, but some rice cultivars are resistant to the nematode such as wild rice *Oryza longistaminata* and *O. glaberrima* which can be used as the parents in breeding resistant varieties (Cabasan et al., 2014).

### Chemical Strategies

Chemical management of *Meloidogyne graminicola* is common in Asia and various methods are used. Seeds can be treated with non-fumigant nematicides such as carbofuran, and seedling roots can be dipped in systemic non-fumigant chemicals such as oxamyl, phorate and carbofuran. Fumigation with chemicals such as 1, 3-dichloropropene can help to reduce the number of nematodes before planting (Dutta et al., 2012).

### Other Strategies

Different biological control agents such as beneficial fungi i.e. *Purpureocillium* sp. *Trichoderma harzianum*, *T. virens*, *Catenaria anguillulae*, and beneficial bacteria i.e. *Bacillus* sp., *Pseudomonas* sp. are potential biological control organisms of *Meloidogyne graminicola* (Pankaj et al., 2015; Mantellin et al., 2017; Bui et al, 2020).

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