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# MELOIDOGYNE SPP. IN MACEDONIA: DISTRIBUTION AND VIRULENCE FOR Mi RESISTANCE IN TOMATO

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**Summary**. The occurrence of *Meloidogyne* in several areas of Macedonia was surveyed. Seventy-three isolates from nine locations (fields and glasshouses) were identified and their behaviour on two susceptible and two *Mi* resistant tomato cultivars compared. *M. incognita* (47,9%) and *M. javanica* (35,6%) are the predominant species followed by *M. arenaria* (13,7%), and, sporadically, *M. hapla* (2,7%) was found. Mixtures of species were present at nearly all locations. Virulent isolates were found in *M. incognita* (11%) and *M. javanica* (46%) as well as in *M. arenaria* (50%). *M. hapla* isolates were compatible with all tomato genotypes tested. The impact of the occurrence of virulence for *Mi* carrying tomato genotypes is discussed.

The widely distributed polyphagous root knot nematodes (Meloidogyne spp.) are among the most damaging plant parasitic nematodes in many economically important crops. Damage by the different species, its populations (intraspecific variation) and subsequent control are important issues. A means of control that is environmentally satisfactory is the growing of resistant cultivars but this is generally hampered because resistance is often not universal i.e. not directed to all Meloidogyne species present in an area or by the occurrence of virulent field populations. Resistance against root knot nematodes has been described for various crops, among them tomato (Lycopersicum esculentum) (e.g. Fassuliotis, 1987; Robert, 1995).

Resistance in tomato to root knot nematodes was found about 50 years ago in an accession of *L. peruvianum* and introgressed in *L. esculentum* using embryo rescue (Smith, 1944). All currently available root-knot resistant tomato

cultivars are derived from this source (Medina-Filho and Tanksley, 1983). A major gene (*Mi*) located on chromosome 6 (Gilbert and McGuire, 1956) controls resistance. The *Mi* gene confers resistance to *M. incognita*, *M. javanica* and *M. arenaria* but not to *M. hapla* (Hadisoeganda and Sasser, 1982; Ammati *et al.*, 1985).

There are two major limiting factors in the use of *Mi*: (i) at high soil temperature (28-30 °C) the resistance does not function (Dropkin, 1969) and (ii) the occurrence of resistance breaking root knot nematode populations. *M. incognita, M. arenaria* and *M. javanica* include virulent field populations and isolates selected on *Mi*-bearing cultivars after several generations (e.g. Riggs and Winstead, 1959; Roberts and Thomason, 1986; Ogallo and McClure, 1996; Molinari and Miacola, 1997). Variation in virulence in *M. incognita* populations towards the *Mi* gene has been studied by many authors (Jarquin-Barberena *et al.*, 1991; Castagnone-Sereno

et al., 1992, 1996; Anwar et al., 1994; Mani and Zidgali, 1995). There are also several reports on virulent M. arenaria and M. javanica populations (Sikora et al., 1973; Prot, 1984; Noe, 1992; Tzortzakakis and Gowen, 1996). Currently seven additional independent dominant Meloidogyne resistance genes, designated Mi-2 to Mi-8, have been identified from L. peruvianum accessions and one from L. chilense. These genes are not yet available in commercial cultivars. They display resistance with different properties from Mi and some are resistant to M. hapla or confer resistance at 32 °C (see: Williamson, 1998). Because L. esculentum and L. peruvianum do not normally cross, the behaviour of these resistance genes in a tomato background is as yet hardly known.

In spite of the economic importance of the tomato crop in Macedonia, knowledge on the distribution of *Meloidogyne* species is scarce and even completely absent if intraspecific variations in virulence are considered. These omissions seriously impede implementation of proper management systems for *Meloidogyne*. As a first step towards such a system we report in this paper on the distribution of *Meloidogyne* populations (a)virulent on *Mi*-bearing tomato genotypes in Macedonian vegetable and tobacco fields.

#### Materials and methods

The susceptible tomato, *L. esculentum* Mill., cvs Moneymaker and Vivia - F 172, and the *Mi* resistant genotypes cv. Carmello GC 204 and cv. Manthos GC 785 were used. S&G Sandoz Seeds, The Netherlands, kindly provided us with the seeds, except for cv. Moneymaker. To exclude nullifying the *Mi* gene at high temperature (Dropkin, 1969) experiments were done at 20-25 °C in a glasshouse. Two week old seedlings were transplanted into 20 cm plastic pots filled with sterilised sand and allowed to establish for two weeks before inoculation with second stage juveniles. Pots were fertilised at regular

intervals and watered with tap water as required. To avoid contamination pots were kept separate on saucers.

In 1996 and 1997 populations of Meloido-

gyne from nine areas (Table I), seven cropped with vegetables (mainly tomatoes) in glasshouses and two tobacco fields were collected from infested roots and rhizosphere soil. Only in the Gevgelija glasshouse was the Mi resistant tomato cv. Suzo grown. The seventy-three isolates (lines) were set up from single egg masses and propagated and maintained for about seven months on cv. Moneymaker. To obtain enough second stage juveniles (J<sub>2</sub>) for the virulence tests eggs were harvested approximately 12 weeks after inoculation by dissolving egg masses in 0.5% NaOCl-solution (Hussey and Barker, 1973). J<sub>2</sub>s were hatched in water and stored at 4 °C until required. The average egg mass contained 200 J<sub>2</sub>s. Prior to inoculation nematodes were allowed to pass through a cotton filter and viable juveniles were collected after one day. In virulence tests 400 J<sub>2</sub>s (~ 2 egg masses (P<sub>i</sub>)) in suspension were pipetted with an automatic syringe onto the soil surface around the stem base of the test plant seedlings followed by light watering. Seven weeks later plants were harvested individually, their roots washed free from sand stained with Phloxine B solution (Daykin and Hussey, 1985) and the egg masses (P<sub>f</sub>) and galls counted. Multiplication rates (Pf/Pi) were assessed. Host status was divided into three categories as follows:  $P_f/P_i \ge 1.0$ , suitable host (SH);  $0.1 < P_f/P_i < 1.0$ , poor host (PH) and  $P_f/P_i \le 0.1$ , non-host (NH). Each nematode population was tested in four replicates on the susceptible and resistant genotypes. For each isolate, after square root transformation of the data (data not shown), pair-wise differences between the means were analysed for significance and Least Significant Difference (LSD) at P=0.05 with ANOVA using Genstat (Payne et al., 1987).

Species composition of the isolates was determined in two ways. (i) Using the morphological criteria of perineal pattern of adult females

TABLE I - Origin, preceding crop and species designation of populations and isolates used.

Location	Location Preceding crop		Number of isolates	Species identified	Code*	
Bogdanci	tomato, cv. Priska	1	7	M. incognita	MiB	
Hamzali	cucumber	3	12	M. incognita	MiSH	
		1	2	M. javanica	MjSH	
	tomato, cv. Balka	1	3	M. arenaria	MaSH	
Bansko	cucumber	1	4	M. incognita	MiSB	
Ilovica	tobacco	2	5	M. incognita	Mil	
Prosenikovo	cucumber	2	7	M. incognita	MiP	
		1	2	M. javanica	MjP	
		1	1	M. arenaria	МаР	
Kocani	cucumber	1	3	M. javanica	MjK	
	tomato, cv. Balka	2	8	M. javanica	MjK	
		1	1	M. arenaria	MaK	
		1	1	M. hapla	MhK	
Gevgelija	tomato, cv. Suzo	1	7	M. javanica	MjG	
Kumanovo	tomato, cv. Vivia	1	4	M. javanica	MjKu	
		1	1	M. hapla	MhKu	
Stuka	tobacco	1	5	M. arenaria	MaS	

<sup>\*</sup> isolates are coded as indicated followed by numbers for field population and isolate.

and morphometrics of males and J2s (Jepson, 1987). At least 15 perineal patterns and 25 males and juveniles of each isolate were examined. (ii) By amplifying the intergenic spacer region between the cytochrome oxidase II gene and the 16S rRNA gene in the mitochondrial genome of single juveniles (Powers and Haris, 1993). In this approach ten individual J<sub>2</sub>s were handpicked, homogenised in a 15 µl drop of sterile water, frozen for future analysis or immediately processed in a PCR procedure in a final volume of 25 µl as described. Following DNA amplification the products were separated on agarose gel, stained with ethidium bromide and visualised on a UV box. M. arenaria is characterised by a unique 1.1 kb fragment. To allow discrimination among species with identically sized amplification products standard restriction digestions (Dra I or Hinf I) of 5 µl of the amplified products were conducted for 2-4 hours at 37 °C and evaluated on 1.5% agarose gels. Di-

gestion with Dra I was carried out in case of the presence of a 0.52 kb amplification product. A four-banded pattern (0.44, 0.29, 0.23 and 0.08 kb fragments) separated *M. bapla* from *M. chitwoodi, M. marylandi, M. nataliei, M. naasi* and *M. fallax.* In case of the presence of a 1.7 kb fragment a Hinf I digestion distinguished *M. javanica* from *M. incognita* on basis of an undigested band and two-banded patterns (0.4 and 1.3 kb fragments) respectively (Williamson *et al.*, 1994). Results were obtained with about 90% of the individual J<sub>2</sub>s tested.

#### Results

There were hardly any discrepancies between the time consuming morphological identifications and the rather straightforward and rapid identifications based on amplification of intergenic spacer region of the mtDNA (Fig. 1). Contradictory results between the morphological and DNA-based identifications were obtained with three isolates (4.1%). The perineal patterns pointed at *M. javanica* and the DNA based identifications at *M. incognita*. The outcomes of the DNA analyses were followed.

Seventy-three isolate originating from 22 *Meloidogyne* populations were collected in 11 areas previously cropped with cucumber, tobacco or tomato. Two glasshouses turned out to be free from *Meloidogyne* infestations. Mostly mixtures of species were found and *M. javanica* (Treub) Chitw. was present in nearly all tomato crops. Thirty-five isolates were identified as *M. incognita* (Kofoid *et* White) Chitw., 26 as *M. javanica*, ten as *M. arenaria* (Neal) Chitw. and two as *M. hapla* Chitw.

Virulence characteristics of the isolates were determined on the susceptible cvs Moneymaker and Vivia and the *Mi* resistant cvs Carmello and Manthos (Table II, Fig. 2). All *M. incognita* isolates reproduced on both susceptible cultivars. It was noticed that often the numbers of egg masses on cv. Moneymaker were higher than the numbers of galls. On cv. Vivia the opposite was true. Four out of the 35 *M. incognita* isolates (11%) (two MiB, MiS and Mil) also produced egg masses on the resistant cultivars. One of the two MiB isolates, being the most virulent, reproduced equally well on cv. Moneymaker and both resistant cultivars.

As also shown in Table II the *M. javanica* isolates generally reproduced well on both susceptible cultivars although numbers of egg

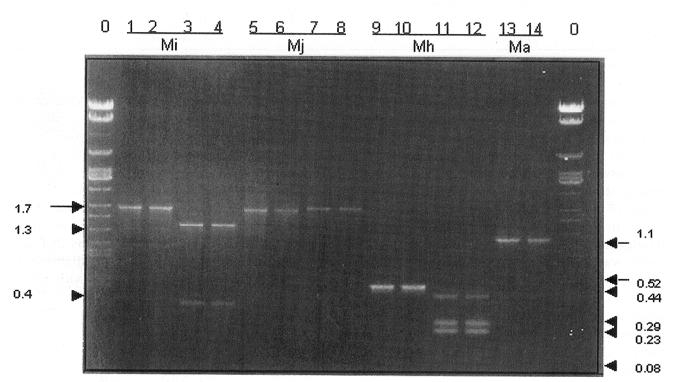


Fig. 1 - PCR amplification of mtDNA of individual J2s. Primers and conditions are as described by Powers and Harris (1993). The 1.7 kb product is characteristic for *Melidogyne incognita* and *M. javanica* (lanes 1, 2 and 5, 6 respectively). After a Hinf I digestion of the 1.7 kb fragment *M. incognita* (lane 3, 4) is identified by a two banded pattern and the *M. javanica* (lane 7, 8) fragments remains undigested. The 0.52 kb product (lanes 9, 10) is characteristic for a number of *Meloidogyne* species. After a Dra I digestion four unique fragments identify *M. bapla* (lanes 11, 12). *M. arenaria* gives a 1.1 kb product (lanes 13, 14). The first and last lanes are standard markers.

Table II - Means of the square root numbers of egg masses of 73 Meloidogyne isolates on roots of four Lycopersicum genotypes differing in presence or absence of the Mi gene (means are of four replicates; virulent populations are shaded).

Isolates	Egg masses / Root system					Egg masses / Root system			
	Money- maker	Vivia	Carmello	Manthos	Isolates	Money- maker	Vivia	Carmello	Manthos
MiB 12.4	8,25 *a	4,47 **b	0,00 с	1,41 c	MiP 33.3	9,33 a	8,25 a	0,00 с	0,00 с
MiB 12.5	10,72 *a	6,40 **b	1,41 c	0,00 d	MiP 33.6	5,74 a	4,00 *b	0,00 c	0,00 c
MiB 12.6	7,35 *a	11,83 *b	0,00 c	0,00 c	MiP 33.8	9,06 a	8,72 a	1,00 c	1,00 c
MiB 12.7	3,46 *a	9,64 **b	2,45 **a	2,65 a	MjP 31.1	9,33 a	5,00 **b	0,00 c	1,00 d
MiB 12.8	8,83 *a	9,43 **a	0,00 c	0,00 с	MjP 31.2	10,39 *a	8,89 **b	0,00 *c	0,00 c
MiB 12.9	9,49 *a	8,43 a	2,00 c	5,10 d	MaP 33.5	8,12 *a	7,07 **a	0,00 c	0,00 c
MiB 12.10	5,66 *a	7,28 **b	1,00 c	0,00 d	MjK 21.1	8,66 a	3,61 b	2,65 bc	2,00 c
MiSH 17.4	6,93 *a	0,00	0,00 с	0,00 с	MjK 21.4	3,61 *a	5,29 *b	1,00 *c	2,45 d
MiSH 17.5	7,87 *a	4,47 **b	0,00 *c	0,00 *c	MjK 21.8	8,89 a	6,00 *b	2,45 c	1,41 d
MiSH 17.7	7,35 a	6,56 a	0,00 c	0,00 c	MjK 2.2	8,25 a	7,00 b	0,00 c	0,00 c
MiSH 17.9	10,15 a	12,04 *b	0,00 c	1,00 d	MjK 2.4	9,59 a	5,83 b	0,00 c	0,00 c
MiSH 17.10	16,09 *a	10,10 *b	1,00 c	0,00 d	MjK 2.6	7,35 a	4,80 b	0,00 c	1,00 d
MiSH 17.12	6,00 a	18,28 *b	0,00 c	1,41 d	MjK 2.7	9,49 a	3,61 **b	0,00 c	0,00 c
MiSH 25.1	6,78 *a	8,77 b	0,00 c	0,00 c	MjK 2.10	11,05 *a	6,86 *b	0,00 c	0,00 c
MiSH 25.2	11,96 a	10,91 *a	1,00 c	0,00 d	MjK 3.3	8,25 *a	8,60 a	1,00 c	3,46 d
MiSH 25.3	8,19 ***	13,27 **b	0,00 **c	1,41 *d	MjK 3.12	7,87 a	4,58 b	0,00 c	1,00 d
MiSH 25.8	13,49 *a	9,80 *b	2,00 c	0,00 d	MjK 3.13	14,35 a	10,10 b	1,00 c	0,00 d
MiSH 26.2	9,90 a	9,43 a	0,00 c	0,00 c	MaK 22.1	11,58 a	12,17 a	2,83 *c	4,58 *d
MiSH 26.3	13,67 a	10,30 **b	0,00 **c	1,00 d	MhK 2.8	8,66 a	6,71 a	1,73 c	3,61 a
MjSH 15.3	8,72 a	2,65 *b	0,00 c	1,00 d	MjG 4.3	9,22 a	6,48 b	1,41 c	3,16 d
MjSH 15.10	8,49 a	3,74 b	2,24 c	1,00 d	MjG 4.4	3,16 a	3,46 a	1,00 c	0,00 d
MaSH 25.4	15,52 a	14,63 *a	1,41 c	1,00 c	MjG 4.5	9,27 a	12,53 a	2,24 c	8,89 a
MaSH 25.5	7,07 a	7,35 a	1,41 c	1,41 c	MjG 4.6	11,40 a	5,20 *b	3,00 c	2,65 c
MaSH 25.10	14,18 a	11,18 **b	1,41 c	2,00 c	MjG 4.10	7,48 *a	2,83 **b	1,41 *c	0,00 d
MiSB 19.3	9,38 *a	8,06 a	1,00 c	1,00 c	MjG 4.11	9,00 a	4,69 b	1,73 c	1,41 c
MiSB 19.6	16,22 a	9,95 b	1,00 *c	0,00 d	MjG 4.13	8,49 a	4,58 b	0,00 c	1,41 d
MiSB 19.7	7,21 a	4,90 *b	1,00 c	0,00 *d	MjKu 7.3	9,59 *a	4,90 *b	1,41 c	2,00 c
MiSB 19.9	9,59 a	8,37 *a	0,00 c	0,00 c	MjKu 7.4	8,37 a	7,21 a	0,00 c	0,00 c
Mil 29.3	3,87 a	2,83 a	0,00 c	0,00 c	MjKu 7.7	10,20 *a	7,35 **b	1,00 c	2,24 d
Mil 29.6	3,16 a	3,16 a	0,00 c	0,00 c	MjKu 7.9	8,72 *a	5,57 *b	0,00 *c	1,73 d
Mil 29.10	11,09 a	8,25 b	1,00 c	2,00 *d	MhKu 7.6	9,90 *a	5,29 **b	2,00 *c	2,65 **b
Mil 30.4	7,21 a	7,35 a	0,00 c	0,00 c	MaS 27.2	8,49 a	8,00 a	0,00 *c	0,00 *c
Mil 30.7	6,93 a	7,68 a	1,41 c	0,00 d	MaS 27.5	13,67 a	12,49 a	0,00 c	1,41 d
MiP 32.5	7,62 a	5,10 *b	0,00 c	0,00 c	MaS 27.6	11,58 a	11,09 a	2,24 c	3,87 d
MiP 32.9	4,00 a	2,00 *b	0,00 c	0,00 c	MaS 27.9	8,83 a	7,94 *a	2,65 *c	2,00 c
MiP 32.11	4,58 a	2,45 b	0,00 c	0,00 c	MaS 27.11	7,28 a	6,16 a	0,00 c	2,00 d
MiP 33.1	6,71 a	5,83 a	0,00 c	0,00 c					

<sup>\*</sup> numbers of galls less that numbers of egg masses. \*\* numbers of galls more that numbers of egg masses. Isolate-genotype combinations sharing a common letter do not differ significantly at P=0.05.

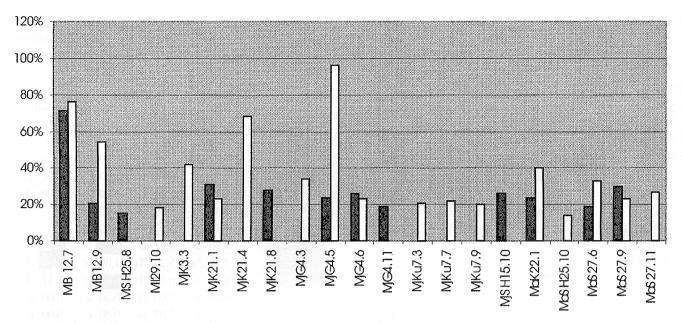


Fig. 2 - Multiplication rates of four *M. incognita*, twelve *M. javanica* and five *M. arenaria* isolates virulent to two *Mi*-bearing *Lycopersicum* genotypes relative to the susceptible cv. Moneymaker. cv. Carmello, cv. Manthos.

masses on cv. Vivia tended to be less than on cv. Moneymaker. Twelve out of the 26 isolates (46%) were virulent at least on one of the *Mi*-bearing cultivars. Some of the isolates (MjG, MjK and MjSH) produced slightly more egg masses on cv. Carmello than on cv. Manthos, while for some other isolates from the same locations (MjG, MjK and MjKu) the opposite was true.

The highest number of virulent isolates (50%) was found with *M. arenaria*. All isolates reproduced on the two susceptible cultivars. Isolate MaK and two MaS isolates were also fully compatible with the both resistant cultivars and MaSH and one of the MaS lines reproduced better on cv. Manthos than on cv. Carmello.

Both *M. hapla* isolates were fully compatible with all four tomato genotypes.

## Discussion

To distinguish *M. hapla* from a set of other *Meloidogyne* spp. a four-banded instead of a two-

banded pattern (Powers and Harris, 1993) was found upon Dra I digestion of the 0.52 kb amplification products and electrophoresis on an agarose gel. The 0.29 kb and 0.23 kb bands were as published. Apparently a second 0.52 kb product was amplified which was digested into a 0.44 kb and a weak 0.08 kb band. Similarly Williamson et al. (1994) found deviating results after a Hinf I digestion of the PCR amplification products and electrophoresis on agarose of M. incognita and M. javanica with a two-banded (1.3 an 0.4 kb) and a single undigested 1.7 kb band respectively. Powers and Harris (1993) published the occurrence of three (1.0, 0.4 and 0.3 kb) and two (1.0 and 0.7 kb) bands (Fig. 1). The reason for such discrepancies is not known but is probably due to differences in the origin of the PCR enzymes used.

In this survey on the occurrence of *Meloido-gyne* in Macedonia 73 isolates originating from nine locations in vegetable and tobacco growing areas were identified and their behaviour on two susceptible and two *Mi* resistant tomato cultivars compared. *M. incognita* (47.9%) and

*M. javanica* (35.6%) predominated followed by *M. arenaria* (13.7%), and, sporadically, *M. hapla* (2.7%) was found. Often species mixtures were found. These figures are consistent with other surveys on *Meloidogyne* in the Mediterranean area (Sorribas and Verdejo-Lucas, 1994; Eddaoudi *et al.*, 1997; Tzortzakakis, 1997).

All isolates reproduced well on both susceptible cultivars. Cv. Moneymaker was in general a better host than cv. Vivia. The virulent *M. incognita*, *M. javanica* and *M. arenaria* isolates reproduced, although to different degrees, on both resistant cultivars (Fig. 2). Most of the isolates reproduced better on cv. Manthos than on cv. Carmello. As expected the two *M. hapla* isolates were fully compatible with the four tomato genotypes (Medina-Filho and Stevens, 1980; Hadisoeganda and Sasser, 1982; Santo, 1982; Ammati *et al.*, 1985; Brown *et al.*, 1997).

Many authors reported on variability in virulence in field populations of M. incognita, M. javanica and M. arenaria from different parts of the world (Roberts et al., 1990; Castagnone-Sereno et al., 1993; Guskova and Al-Bakkur, 1996; Anwar et al., 1996; Kaloshian et al., 1996). Virulent populations may occur naturally or arise from avirulent populations after repeated selection pressure by Mi-carrying cultivars (e.g. Castagnone Sereno et al., 1993). All tomato glasshouses surveyed were, as far as known, until now cropped with susceptible cultivars with the exception of Gevgelija where the Mi resistant cv. Suzo was grown. Tzortzakakis and Gowen (1996) found fully virulent M. javanica lines from populations never exposed to the Mi gene. Therefore it remains unclear if the fairly high incidence of virulent M. javanica isolates from the Gevgelija area resulted from selection or that they are already present naturally.

In Macedonia, glasshouses for tomato cropping are normally treated with systemic nematicides or disinfected by steam sterilisation in December. Tomatoes are transplanted in January. After harvest early in the summer glasshouses are left fallow and often in the autumn a second

different crop will be grown. The results presented here show that growing *Mi*-resistant tomato cultivars in a cropping system as outlined will be beneficial in case virulent populations are present. The set backs are mainly found in damage to a second crop in case high *Meloidogyne* densities have built up in the spring. This problem is even more pronounced in tomatoes grown as a field crop because of the absence of soil disinfection and the relatively high numbers of (a)virulent *Meloidogyne* populations found.

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