EPPO Datasheet: Meloidogyne fallax

Last updated: 2020-10-06

IDENTITY

Preferred name: Meloidogyne fallax

Authority: Karssen

Taxonomic position: Animalia: Nematoda: Chromadorea:

Rhabditida: Meloidogynidae

Common names: false Columbia root-knot nematode

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EU Categorization: A2 Quarantine pest (Annex II B)

EPPO Code: MELGFA



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Notes on taxonomy and nomenclature

Meloidogyne fallax was detected for the first time in 1992 in a field plot experiment 1.5 km north of Baexem (NL), and was initially considered as a deviant *M. chitwoodi* Golden population (Karssen, 1994). On the basis of differences in isozyme patterns, *M. fallax* was proposed as a new race of *M. chitwoodi* (van Meggelen *et al.*, 1994), and named *M. chitwoodi* B-type (Karssen, 1995). As more differences between *M. chitwoodi* and the B-types were discovered, this race status became unacceptable, and *M. fallax* was described as a new species (Karssen, 1996).

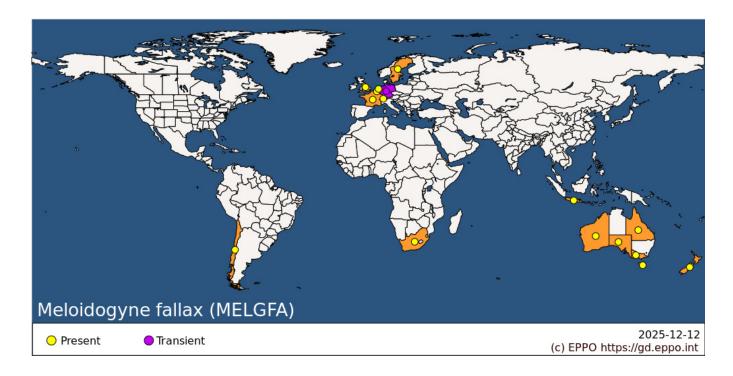
HOSTS

M. fallax was detected on and described from potato (Solanum tuberosum). Host-range includes a wide range of dicotyledonous and monocotyledons plants, including weeds, ornamentals, and economically important crops such as carrot (Daucus carota), black salsify (Scorzonera hispanica) and tomato (Solanum lycopersicum). The experimental host range of M. fallax mostly overlaps that of M. chitwoodi, but differential hosts have been found. For example dwarf beans (Phaseolus vulgaris), valerian (Valeriana officinalis), maize (Zea mays), Erica cinerea and Potentilla fruticosa are good hosts for M. chitwoodi and not for M. fallax, while the reverse is the case for Oenothera glazioviana, Phacelia tanacetifolia, Hemerocallis cv. Rajah and Dicentra spectabilis (Brinkman et al., 1996). It is expected that many more plant species will be hosts of M. fallax than currently known, since this is the case also with other, closely related root knot nematodes.

Host list: Allium porrum, Asparagus officinalis, Avena strigosa, Beta vulgaris, Cichorium intybus, Cynara scolymus, Daucus carota subsp. sativus, Fragaria x ananassa, Hemerocallis sp., Hordeum vulgare, Lactuca sativa, Lamprocapnos spectabilis, Leptinella sp., Lolium multiflorum, Medicago sativa, Oenothera glazioviana, Phacelia tanacetifolia, Scorzonera hispanica, Solanum lycopersicum, Solanum nigrum, Solanum physalifolium, Solanum tuberosum, Trifolium repens, Triticum aestivum subsp. aestivum

GEOGRAPHICAL DISTRIBUTION

After the first record near Baexem (NL) in 1992, *M. fallax* was recorded on potato at several locations in the southern and south-eastern part of the Netherlands (Karssen, 1996), close to the German and Belgium borders. Within the EPPO region it was detected locally in Belgium, France, Germany, Sweden, Switzerland and the United Kingdom (England). In addition, Topalovi? *et al.* (2017) revealed that a *Meloidogyne* species detected in Ireland in 1965 belongs to *M. fallax*. *M. fallax* has never been reported from the natural environment in Europe. Outside Europe it has been reported from Australia, Chile, New Zealand and South Africa. New Zealand is the only known country where *M. fallax* is widely distributed (North and South Island) and detected in cropping and pasture fields (Rohan *et al.*, 2016), strongly suggesting that it could be the place of origin of this pest.



EPPO Region: Belgium, France (mainland), Germany, Netherlands, Sweden, Switzerland, United Kingdom

(England)

Africa: South Africa Asia: Indonesia (Java) South America: Chile

Oceania: Australia (Queensland, South Australia, Tasmania, Victoria, Western Australia), New Zealand

BIOLOGY

The life cycles of *M. fallax* and *M. chitwoodi* are, in general, the same with respect to root penetration, gall induction, symptomatology, number of moults, parthenogenetic reproduction and chromosome number: Both *M. chitwoodi* and *M. fallax* usually reproduce by parthenogenesis. They can have one to three generations per year in the Netherlands and produce several hundreds of eggs per female, deposited in an egg sac. These egg sacs allow the eggs to survive under unfavorable conditions (EFSA, 2019). Initial results by van der Beek (1997) indicated that *M. fallax* had a shorter life cycle than *M. chitwoodi* in a virulence study on potato.

Host races, as described for *M. chitwoodi*, have not been detected for *M. fallax* so far. Successful hybridization was not obtained when *M. fallax* and *M. chitwoodi* were crossed in two different experiments; the F1 was viable, but the F2 second-stage juveniles were not viable and showed morphological distortions (van der Beek & Karssen, 1997).

In addition, differences in terms of hatching responses to root diffusate and host age between *M. chitwoodi* and *M. fallax* suggests different survival strategies between these two species (Wesemael *et al.*, 2006).

M. fallax, M. chitwoodi and M. minor Karssen are closely related, morphologically and at DNA level. Phylogenetically they appear to be in the same distinct clade within the genus Meloidogyne (Holterman et al., 2009; Elling, 2013).

DETECTION AND IDENTIFICATION

Symptoms

Above-ground symptoms of heavily infested plants include stunting and yellowing, while below ground galling is typical. The root galls produced by *M. chitwoodi* and *M. fallax* are comparable to those produced by several other

root-knot species, i.e. relatively small galls in general without secondary roots emerging from them (these secondary roots are seen in *M. hapla*). On potato tubers, *M. chitwoodi* and *M. fallax* cause numerous small pimple-like raised areas on the surface (in *M. hapla* these swellings are not evident). Some potato cultivars, although heavily infested, may be free from visible external symptoms, while the internal potato tissue is necrotic and brownish, just below the peel (EPPO, 2016a).

Morphology

Sedentary females are annulated, pearly white and globular to pear-shaped, $400-720~\mu m$ long and $250-460~\mu m$ wide. The stylet is dorsally curved, $13.9-15.2~\mu m$ long, with rounded to ovoid stylet knobs, slightly sloping posteriorly. The non-sedentary males are vermiform, annulated, slightly tapering anteriorly, bluntly rounded posteriorly, $735-1520~\mu m$ long and $27-44~\mu m$ wide. The stylet is $18.9-20.9~\mu m$ long, with large rounded knobs, set off from the shaft. The non-sedentary second-stage juveniles are vermiform, annulated, tapering at both ends, $380-435~\mu m$ long, $13.3-16.4~\mu m$ wide, with a tail length of $46-56~\mu m$ and a hyaline tail part $12.2-15.8~\mu m$ in length (Karssen, 1996).

M. fallax is closely related morphologically to *M. chitwoodi*, and this misleading resemblance was the reason for giving the species its name. The most striking differences for males and females are stylet length (longer for *M. fallax*) and stylet knob shape (*M. fallax*: prominent and rounded; *M. chitwoodi*: small and irregular). The second-stage juveniles differ in mean body length, tail length and hyaline tail length (all longer for *M. fallax*). With the scanning electron microscope, it can be observed that the male head of *M. fallax* has an elevated labial disk. Differences exist in the female perineal pattern (*M. fallax* has a relatively higher dorsal arch and thicker striae) (EPPO, 2016a; see also Karssen, 2002).

The species can be reliably distinguished by morphological observation of females, males, and second-stage juveniles in combination with biochemical (isozyme electrophoresis) or molecular (PCR) methods; see EPPO, 2016. To predict the amount of potato tuber damage caused by *M. fallax*, a quantitative DNA-test was developed for soil (Hay *et al.*, 2016).

Detection and inspection methods

Specific guidance on the sampling of soil and potato tubers is given in the EPPO Standards PM 9/17 (EPPO, 2013b), PM 3/71 (EPPO, 2007) and PM 3/69 (EPPO, 2019a). Populations in the soil rapidly decline in the absence of a host and nematodes reproduce better on a good host. Therefore, detection of the nematodes through field inspection and soil sampling is more sensitive if done as close as possible to the time of harvest of a host crop, targeting particularly susceptible plants (EPPO, 2013b). In each lot of potato (typically 25 tonnes), 200 tubers are randomly sampled and processed (EPPO, 2019a).

Nematode extraction and identification should be carried out according to EPPO Standards PM 7/119 (EPPO, 2013a) and PM 7/41 *Diagnostic protocol for* Meloidogyne chitwoodi *and* M. fallax (EPPO, 2016a).

Historically, in order to distinguish M. fallax from other Meloidogyne spp., molecular methods have been used. Karssen et al. (1995) discriminated M. fallax and M. chitwoodi females by their esterase (EC 3.1.1.1) and malate dehydrogenase (EC 1.1.1.37) isozyme patterns, using the general method of Esbenshade & Triantaphyllou (1985) for identification of female Meloidogyne species by isozyme electrophoresis. Additionally, the isozyme glucose 6phosphate dehydrogenase (EC 1.1.1.49) was used to differentiate the two species (van der Beek & Karssen, 1997). van der Beek et al. (1997) and Tastet et al. (1999) used mini two-dimensional gel electrophoresis to study the total soluble protein patterns of M. hapla, M. chitwoodi and M. fallax, and confirmed these species to be distinct biological groups. An overwhelming number of molecular tests has been developed to identify M. fallax from soil, root or tubers and to separate it from related species such as M. chitwoodi, including PCR (Peterson & Vrain, 1996, Peterson et al., 1997; Wishart et al., 2002), PCR RFLP (Zijlstra et al. 1995, Zijlstra, 1997) AFLP (v.d. Beek et al., 1998; Fargette et al., 2005), SCAR (Zijlstra, 2000; Fourie et al., 2001), real time TaqMan PCR (Zijlstra & v. Hoof, 2006; de Haan et al., 2014), RAPD (Adam et al., 2007), satDNA (Castagnone-Sereno et al., 1999; Castagnone-Sereno, 2000; Mestrovich et al., 2009 & 2013), LAMP (Zhang & Gleason, 2019), HRMC (Holterman et al., 2012), barcoding (Hodgetts et al., 2016, EPPO, 2016b). A serological test was also developed to separate M. fallax from M. chitwoodi and other root-knot nematodes (Tastet et al., 2001). Only a selection of the above-mentioned tests are recommended for identification (see EPPO, 2016a).

PATHWAYS FOR MOVEMENT

M. fallax has very limited potential for natural movement; only second-stage juveniles can move in the soil and, at most, only a few tens of centimetres. The most likely pathway for introducing M. fallax into a new area is through the movement of infested or contaminated planting material. Infested host plants or host products such as bulbs or tubers can easily transport the nematode. The movement of non-host plants for planting (e.g. seedling transplants, nursery stock), non-host plant products (e.g. bulbs, tubers, corms and rhizomes), equipment and machinery which are contaminated with soil infested with M. fallax could also result in spread (EPPO, 2013b). Soil as such is also a possible pathway. Infective juveniles of this genus have been known to persist for more than one year in the absence of host plants. Nematode movement can also be facilitated by contaminated irrigation water.

PEST SIGNIFICANCE

Economic impact

In trials, *M. fallax* caused the same symptoms on potato tubers, black salsify and carrots as *M. chitwoodi*, i.e. external galling and internal necrosis just below the skin (Brinkman *et al.*, 1996; van Riel & Goossens, 1996). The reported natural outbreaks of *M. fallax* on potato showed these external symptoms (Karssen, 1996). Goosens (1995) reported infected *Asparagus officinalis* and several ornamentals with root-knots in an experimental field with an infestation of *M. fallax*. *M. fallax* sometimes occurs in mixed infestations with *M. chitwoodi* (Wesemael *et al.*, 2006).

Meloidogyne fallax mainly induces quality losses caused by cosmetic damage on black salsify, leek, carrot and potato. If the level of infection is high a complete rejection of these crops is possible. So far, this nematode has a limited damaging effect on other known host crops (EFSA, 2012).

In countries where *M. fallax* is regulated, traded plants and plant products infested with *M. fallax* may need to be destroyed.

Control

There is no direct practical experience of the control of *M. fallax*. Research on *M. fallax* in the Netherlands has focused on host suitability, damage thresholds, effect of fallow, the use of green manure crops and time of sowing. The first results indicate that fallow for one year reduced the population by more than 95%, but this reduction was not sufficient to ensure that subsequent crops met quality standards. There was less damage in sugar beet and carrot when these crops were sown later in spring. Farmers are advised to be careful when growing green-manure crops on infested fields, because some species are suitable hosts for *M. fallax*. *Phaseolus vulgaris* was the only tested crop with no reproduction of *M. fallax*, while maize and cereals were poor hosts (Brommer, 1996). Several green manures were tested, and no reproduction of *M. fallax* was found on *Eruca Sativa* cv Trio, *Tagetus patula* and *Borago* spp., while *Raphanus sativus* was a poor host. *Avena strigosa* turned out to be a very good host (Visser & Molendijk, 2015).

Janssen et al. (1996) tested several wild tuber-bearing Solanum species, to determine the level of resistance to M. hapla, M. chitwoodi and M. fallax. High resistance to M. chitwoodi and M. fallax was observed in genotypes of S. bulbocastanum, S. hougasii, S. cardiophyllum, S. fendleri and S. brachistotrichum. Differential resistance between M. chitwoodi and M. fallax was observed in S. chacoense, S. stoloniferum and S. gourlayi. Resistance to M. fallax was also found in Solanum sparsipilum (Kouassi et al., 2004) and the resistance gene was studied (Kouassi et al., 2006). By crossing wild tuber bearing Solanum species with Solanum tuberosum, successful introgression of resistance to M. fallax and M. chitwoodi into potato was obtained (Janssen et al., 1997).

In addition, useful resistance against *M. fallax* was found in sea beet (*Beta maritima*) (Yu *et al.*, 1999) and used to develop a resistant sugar beet line (*Beta vulgaris*) (Yu, 2001).

Wesemael & Moens (2012) tested ten common bean (*Phaseolus vulgaris*) cultivars and noted that all these cultivars were non-hosts or poor hosts for *M. fallax*.

An interesting potential control method consists of using the hyper-parasitic bacterium *Pasteuria* spp. So far three different species of *Pasteuria* are reported to be able to parasitize *M. fallax*: *P. penetrans*, *P. nishizawae* and *P. hartismeri* (Wishart *et al.*, 2004; Bishop *et al.*, 2007).

Phytosanitary risk

Because *M. fallax* occurs in crops and situations which are similar to those for *M. chitwoodi*, and the two species are closely related and difficult to distinguish, *M. fallax* presents a phytosanitary risk similar to that of *M. chitwoodi* (EFSA, 2012). *M. fallax* may be able to establish in a large proportion of its host area but may only cause significant damage in certain areas and under certain conditions, causing complete crop rejection (e.g. on potato, carrot and/or black salsify). Soils with a coarse texture such as sandy and sandy-loam soils have a higher probability of being contaminated. Narrow rotation or rotation with alternative hosts facilitates a rapid build-up of population levels and therefore increases the risk for establishment of *M. chitwoodi* and *M. fallax* in a particular field. As observed since 1992, *M. fallax* is expected to spread slowly (EFSA, 2012).

PHYTOSANITARY MEASURES

Measures similar to those for other root-knot nematodes would appear relevant, i.e. that rooted host plants for planting (with or without soil), non-host plants for planting with soil attached and plant products with soil attached come from a pest free area, a pest free place of production or are produced under protected cultivation. Alternatively, soil from non-host plants for planting or plant products can be removed. Soil as such can originate from a pest free area or a pest free place of production. Used machineries, equipment, vehicles, and passengers' shoes can be cleaned. Publicity would allow to enhance public awareness on *M. fallax* when travelling.

Specific requirements are recommended in EPPO Standard PM 8/1 Commodity-specific phytosanitary measures for Potato (EPPO, 2017) for seed and ware potatoes to be imported from third countries. In this Standard, seed potato freedom from M. fallax can also be guaranteed by testing the seed potatoes after harvest following EPPO Standard PM 3/69 Meloidogyne chitwoodi and M. fallax: sampling potato tubers for detection (EPPO, 2019a). Ware potatoes freedom can also be guaranteed by implementing EPPO Standard PM 9/17 National regulatory control system for Meloidogyne chitwoodi and M. fallax (EPPO, 2013b). EPPO Standard PM 3/61 details conditions for establishing pest-free areas and pest-free production and distribution systems for quarantine pests of potato (EPPO, 2019b).

Measures to contain or eradicate *M. chitwoodi* and *M. fallax* are described in the national regulatory control system (EPPO, 2013b).

REFERENCES

Adam MAM, Phillips MS & Blok VC (2007) Molecular diagnostic key for identification of single juveniles of seven common and economically important species of root-knot nematode (*Meloidogyne* spp.). *Plant Pathology* **56**, 190-197.

Bishop AH, Gowen SR, Pembroke B & Trotter JR (2007) Morphological and molecular characteristics of a new species of *Pasteuria* parasitic on *Meloidogyne ardenensis*. *Journal of Invertebrate Pathology* **96**, 28-33.

Brinkman H, Goossens JJM & van Riel HR (1996) Comparative host suitability of selected crop plants to *Meloidogyne chitwoodi* and *M. fallax. Anzeiger für Schädlingskunde, Planzenschutz, Umweltschutz***69**, 127-129.

Brommer E (1996) [Control of the root-knot nematode *Meloidogyne fallax*]. *Publicatie Proefstation voor de Akkerbouw en de Groenteteelt in de Vollegrond* no. 81B, pp. 159-163 (in Dutch).

Castagnone-Sereno P, Leroy F, Bongiovanni M, Zijlstra C & Abad P (1999) Specific diagnosis of the two root-knot nematodes, *Meloidogyne chitwoodi* and *M. fallax. Phytopathology* **89**, 380-384.

Castagnone-Sereno P (2000) Use of satellite DNA for specific diagnosis of the quarantine root-knot nematodes *Meloidogyne chitwoodi* and *M. fallax. EPPO Bulletin* **30**, 581-584.

de Haan EG, Dekker CCEM, Tameling WIL, den Nijs LJMF, Bovenkamp GW & van den Kooman-Gersmann M (2014) The MeloTuber Test: a real-time TaqManReq. PCR-based assay to detect the rootknot nematodes *Meloidogyne chitwoodi* and *M. fallax. Bulletin OEPP/EPPO Bulletin* **44**, 166-175.

EFSA (2012) Macleod A, Anderson H, Follak S, van der Gaag DJ, Potting R, Pruvost O, Smith J, Steffek R, Vloutoglou I, Holt J, Karadjova O, Kehlenback H, Labonne G, Reynaud P, Viaene N, Anthoine G, Holeva M, Hostachy B, Ilieva Z, Karssen G, Krumov V, Limon P, Meffert J, Niere B, Petrova E, Peyre J, Pfeilstetter E, Roelofs W, Rothlisberger F, Sauvion N, Schenck N, Schrader G, Schroeder T, Steinmöller S, Tjou-Tam-Sin L, Ventsislavov V, Verhoeven K & Wesemael W. Pest risk assessment for the European Community plant helath: a comparative approach with case studies. Cases: *Meloidogyne chitwoodi* and *M. fallax. EFSA Supporting publications 2012* EN-319. 1053 pp.

EFSA (2019) den Nijs L, Camilleri M, Diakaki M, Schenk M & Vos S. Pest survey card on *Meloidogyne chitwoodi* and *Meloidogyne fallax*. *EFSA supporting publication 2019* **EN-1572**. 20 pp.

Elling AA (2013) Major emerging problems with minor Meloidogyne species. Phytopathology 103, 1092-1102.

EPPO (2007) EPPO Standard PM 3/71(1) General crop inspection procedure for potatoes + corrigendum. *EPPO Bulletin* **37**, 592-597. Available at: https://gd.eppo.int/standards/PM3/

EPPO (2013a) EPPO Standard PM 7/119 (1) Nematode extraction. *EPPO Bulletin* **43**, 471-495. Available at: https://gd.eppo.int/standards/PM7/

EPPO (2013b). EPPO Standard PM 9/17 (1) *Meloidogyne chitwoodi* and *M. fallax. EPPO Bulletin* **43**, 527-533. Available at: https://gd.eppo.int/standards/PM9/

EPPO (2016a) EPPO Standard PM 7/41 (3). *Meloidogyne chitwoodi* and *M. fallax. EPPO Bulletin* **46**, 171-189. Available at: https://gd.eppo.int/standards/PM7/

EPPO (2016b) EPPO Standard PM 7/129(1) DNA barcoding as an identification tool for a number of regulated pests. *EPPO Bulletin* **46**, 501-537. Available at: https://gd.eppo.int/standards/PM7/

EPPO (2019a) EPPO Standard PM 3/69(2) *Meloidogyne chitwoodi* and *M. fallax*: sampling potato tubers for detection. *EPPO Bulletin* **49**, 486-487. Available at: https://gd.eppo.int/standards/PM3/

EPPO (2019b) EPPO Standard PM 3/61(2) Pest-free areas and pest-free production and distribution systems for quarantine pests of potato. *EPPO Bulletin* **49**, 480-481. Available at: https://gd.eppo.int/standards/PM3/

Esbenshade PR & Triantaphyllou AC (1985) Use of enzyme phenotypes for identification of *Meloidogyne* species. *Journal of Nematology* **17**, 6-20.

Fargette M, Lollier V, Phillips M, Blok V & Erutos R (2005) AFLP analysis of the genetic diversity of *Meloidogyne chitwoodi* and *M. fallax*, major agricultural pests. *Comtes Rendus Biologies* **328**, 455-462.

Fourie H, Zijlstra C & McDonald AH (2001) Identification of root-knot nematode species occurring in South Africa using SCAR-PCR technique. *Nematology* **3**, 675-680.

Goossens JJM (1995) Host range test of *Meloidogyne* n.sp. In: Annual Report 1994 Diagnostic Centre, pp. 95-97. Plant Protection Service, Wageningen (NL).

Hay FS, Ophel-Keller K, Hartley DM & Pethybridge SJ (2016) Prediction of potato tuber damage by root-knot nematodes using quantitative DNA assay of soil. *Plant Disease* **100**, 592-600.

Hodgetts J, Ostoja-Starzewski JC, Prior T, Lawson R, Hall J & Boonham N (2016) DNA barcoding for biosecurity: case studies from the UK plant protection program. *Genome* **59**, 1033-1048.

Holterman MHM, Karssen G, van den Elsen SJJ, van Megen HHB, Bakker J & Helder H (2009) SSU rDNA-based

phylogeny of the Tylenchida sheds light on the evolution of plant feeding & establishes relationships among high impact plant parasitic nematodes. *Phytopathology* **99**, 227-235.

Holterman MHN, Oggenfuss M, Frey JE & Kiewnick S (2012) Evaluation of high-resolution melting curve analysis as a new tool for root-knot nematode diagnostics. *Journal of Phytophatology* **160**, 59-66. Janssen GJW, van Norel A, Verkerk-Bakker B & Janssen R (1996) Resistance to *Meloidogyne chitwoodi*, *M. fallax* and *M. hapla* in wild tuberbearing *Solanum* spp. *Euphytica* **92**, 287-294.

Janssen GJW, Janssen R, Van Norel A, Verkerk-Bakker B & Hoogendoorn J (1996) Expression of resistance to the root-knot nematodes, *Meloidogyne hapla* and *M. fallax*, in wild*Solanum* spp. under field conditions. *European Journal of Plant Pathology* **102**, 859-865.

Janssen GJW, van Norel A, Verkerk-Bakker B & Janssen R (1997) Intra- and interspecific variation of root-knot nematodes, *Meloidogyne* spp., with regard to resistance in wild tuber bearing *Solanum* species. *Fundamental and Applied Nematology* **20**, 449-457.

Karssen G (1994) The use of isozyme phenotypes for the identification of root-knot nematodes (*Meloidogyne* species). In: *Annual Report 1992 Diagnostic Centre*, pp. 85-88. Plant Protection Service, Wageningen (NL).

Karssen G (1995) Morphological and biochemical differentiation in *Meloidogyne chitwoodi* populations in the Netherlands. *Nematologica* **41**, 314-315.

Karssen G (1996) Description of *Meloidogyne fallax* n.sp. (Nematoda: Heteroderidae), a root-knot nematode from the Netherlands. *Fundamental and applied Nematology* **19**, 593-599.

Karssen G, van Hoenselaar T, Verkerk-Bakker B & Janssen R (1995) Species identification of cyst and root-knot nematodes frompotato by electrophoresis of individual females. *Electrophoresis* **16**, 105-109.

Karssen G (2002) *The plant-parasitic nematode genus* Meloidogyne *Göldi, 1892 in Europe*. Brill Academic Publishers, the Netherlands. 157pp.

Kouassi AB, Kerlan MC, Sobczak M, Bantec JP, Rouaux C, Ellisseche D & Mugniery D (2004) Resistance to the root-knot nematode *Meloidogyne fallax* in *Solanum sparsipilum*. *Nematology* **6**, 389-400.

Kouassi AB, Kerlan MC, Caromel B, Dantec JP, Fouville D, Manzanares-Dauleux M, Ellisseche D & Mugniery D (2006) A major gene mapped on chromosome XII is the main factor of a quantitatively inherited resistance to *Meloidogyne fallax* in *Solanum sparsipilum*. *Theoritical and Applied Genetics* **112**, 699-707.

Mestrovic N, Phlol M & Castagnone-Serone P (2009) Relevance of satellite DNA genomic distribution in phylogenetic analysis: a case study with root-knot nematodes of the genus *Meloidogyne*. *Molecular Phylogenetics and Evolution* **50**, 204-208.

Mestrovic N, Pavlek M, Car A, Castagnone-Serone P, Abad, P & Plohl M (2013) Conserved DNA motifs, including CENP-B box-like, are possible promotors of satellite DNA array rearrangesments in nematodes. *PloS ONE* **8**, e67328.

Peterson DJ & Vrain TC (1996) Rapid identification of *Meloidogyne chitwoodi*, *M. hapla* and *M. fallax* using PCR primers to amplify their ribosomal intergenic spacer. *Fundamental and applied Nematology* **19**, 601-605.

Peterson DJ, Zijlstra C, Wishart J, Blok V & Vrain T (1997) Specific probes efficiently distinguish root-knot nematode species using signature in the ribosomal intergenic spacer. *Fundamental and applied Nematology* **20**, 619-626.

Rohan TC, Aalders LT, Bell NL & Shah FA (2016) First report of *Meloidogyne fallax* hosted by *Trifolium repens* (white clover): implications for pasture and crop rotations in New Zealand. *Australasian Plant Disease Notes*. **11**, 14.

Tastet, C, Val F, Lesage M, Renault L, Marche L, Bossis, M & Mugniery D (2001) Application of a putative fatty-acid binding protein to discriminate serologically the two European quarantine root-knot nematodes, *Meloidogyne chitwoodi*

and M. fallax from other Meloidogyne species. European Journal of Plant Pathology 107, 821-832.

Topalovi? O, Moore JF, Janssen T, Bert W & Karssen G (2017) An early record of *Meloidogyne fallax* from Ireland. *ZooKeys* **643**, 33–52. https://doi.org/10.3897/zookeys.643.11266

van der Beek JG (1997) Isolate-by-cultivar interaction in *M. hapla, M. chitwoodi* and *M. fallax* on potato. In: *Interaction between root-knot nematodes and Solanum spp. variation in pathogenicity, cytology, proteins and DNA.* pp. 41-53. Thesis, Wageningen Agricultural University, Wageningen (NL).

van der Beek JG & Karssen G (1997) Interspecific hybridization of meiotic parthenogenetic *Meloidogyne chitwoodi* and *M. fallax. Phytopathology* **87,**1061-1066.

van der Beek JG, Folkertsma R, Poleij LM, van Koert PHG & Bakker J (1997) Molecular evidence that *Meloidogyne hapla, M. chitwoodi* and *M. fallax* are distinct biological entities. *Fundamental and applied Nematology* **20**, 513-520.

van der Beek JG, Folkertsma R, Zijlstra C, van Koert PHG, Poleij LM & Bakker J (1998) Genetic variation among parthenogenetic *Meloidogyne* species revealed by AFLPs and 2D-protein electrophoresis contrasted to morphology. *Fundamental and Applied Nematology* **21**, 401-411.

van Meggelen JC, Karssen G, Janssen GJW, Verkerk B & Janssen R (1994) A new race of *Meloidogyne chitwoodi?*. *Fundamental and appliedNematology* **17**,93.

van Riel HR & Goossens JJM (1996) Response of *Meloidogyne fallax* to differential hosts of *M. chitwoodi*. In: *Annual Report 1995 Diagnostic Centre*, pp. 100-101. Plant Protection Service, Wageningen (NL).

Visser JHM & Molendijk LPG (2015) [Host suitability new green manures for plant-parasitic nematodes]. *Publicatie WUR, Praktijkonderzoek Plant & Omgeving* no. 633, 29pp. (in Dutch).

Wesemael WML, Perry RN & Moens M (2006) The influence of the root diffusate and host age on hatching of the root-knot nematodes. *Nematology* **8**, 895-902.

Wesemael WML & Moens M (2012) Screening of common bean (*Phaseolus vulgaris*) for resistance against temperate root-knot nematodes (*Meloidogyne* spp.). *Pest Management Science* **68**, 702-708.

Wishart J, Phillips MS & Blok VC (2002) Ribosomal intergenic spacer: a polymerase chain reaction diagnostic for *Meloidogyne chitwoodi*, *M. fallax* and *M. hapla. Phytophatology* **92**, 884-892.

Wishart J, Blok VC, Phillips MS & Davies KG (2004) *Pasteuria penetrans* and *P. nishizawae* attachment to *Meloidogyne chitwoodi*, *M. fallax* and *M. hapla. Nematology* **6**, 507-510.

Yu M, Heijbroek W & Pakish LM (1999) The sea beet source of resistance to multiple species of root-knot nematode. *Euphytica* **108**, 151-155.

Yu MH (2001) Registration of M6-1 root-knot nematode resistant sugarbeet germplasm. Crop Science 41, 278-279.

Zhang L & Gleason C (2019) Loop-mediated isothermal amplification for the diagnostic detection of *Meloidogyne chitwoodi* and *M. fallax. Plant Disease* **103**, 12-18.

Zijlstra C (1997) A fast PCR assay to identify *Meloidogyne hapla*, *M. chitwoodi* and *M. fallax*, and to sensitively differentiate them from each other and from *M. incognita* in mixtures. *Fundamental and applied Nematology* **20**, 505-511.

Zijlstra C, Lever AEM, Uenk BJ & van Silfhout CH (1995) Differences between ITS regions of isolates of root-knot nematodes *Meloidogyne hapla* and *M. chitwoodi*. *Phytopathology* **85**, 1231-1237.

Zijlstra C (2000) Reliable identification of quarantine root-knot nematodes *Meloidogyne chitwoodi* and *M. fallax* by PCR-based techniques. *EPPO Bulletin* **30**, 575-579.

Zijlstra C & van Hoof RA (2006) A multiplex real-time polymerase chain reaction (TaqMan) assay for the simultaneous detection of *Meloidogyne chitwoodi* and *M. fallax. Phytopathology* **96**, 1255-1262.

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