



NEMATODES AND FUNGI ASSOCIATED WITH APPLE REPLANT DISORDER IN SAMPLED NEW YORK STATE ORCHARDS

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ABSTRACT

The factors associated with unthrifty growth of apple are not fully determined, resulting in inadequate diagnosis of apple replant disorder (ARD). The present study isolated and identified nematodes and fungi from ARD-infested soils to determine factors that interacted to cause ARD. Nematodes were extracted and identified in several orchard soil subsamples. The remaining soils were mixed to give composite soil. Transformed and non-transformed apple host seedlings were planted in the composite soil in a greenhouse, harvested after 10 weeks, and processed for isolation and identification of fungi. *Pratylenchus* and *Xiphinema* nematodes were present in the soils. Root-lesion nematode (RLN) frequency was always 100%, and dagger nematode (DN) frequency ranged from 40-70%. The RLN and DN ranged from 56-350 and 5-58 per 100 cm³ of soil, respectively. Isolation frequency of *Pythium* species was highest (29%), followed by 10% for *Pythium cryptogea*, 4% for *Pythium cambivora* and *Pythium cactorum*, 3% for *Pythium megasperma*, and 2% for *Phytophthora* species. Both transformed and non-transformed apple hosts were infested by at least several fungi. Of the six apple hosts tested, *Pythium* sp. and *F. oxysporum* colonized six each, *Pythium cryptogea*, *C. lucidum* and *C. destructans* colonized three each, *Pythium cactorum*, *Pythium cambivora* and *Pythium megasperma* two each, and *Phytophthora* sp., *F. solani*, *F. equiseti*, *F. acuminatum* and *Rhizoctonia* sp. one each. Consequently, the ARD symptoms observed on diverse apple hosts were associated with combined effects of RLN, DN and various species of *Pythium*, *Phytophthora*, *Cylindrocarpon*, *Fusarium* and *Rhizoctonia*. Whenever unthrifty growth is observed in soil especially that under perennial fruits and continuously cultivated annual crops, similar diagnostic isolation of multiple pathogens should be performed. In developing management strategies, germplasm for the crop in question should be evaluated in multiple sites or in composite soil in order to identify germplasm with broadspectrum resistance or tolerance to the pathogens.

KEY WORDS: Apple Replant Disease, *Malus* sp., *Pratylenchus*, *Xiphinema*, Transgenics

INTRODUCTION

Replant disorders refer to the poor growth of replanted perennial and annual crops and occur worldwide in fields, orchards and nurseries previously cropped or in fresh soils as intrinsic disorders termed "soil sickness" (Sewell, 1981). Apple replant disorders (ARD) cause widespread and serious difficulty in establishment of replanted apple trees (Hoestra, 1968; Utkhede and Smith, 1994). Symptoms of ARD appear as stunting, diminished root hair growth, root necrosis, sparse, browning and decaying fibrous roots, foliar nutrient deficiency, reduced productivity, and in the most severe cases, tree death within the first three years of planting (Jaffee *et al.*, 1982b; Utkhede and Smith, 1992). Death of feeder roots account for impaired uptake of water and nutrients, whereas damage to root hairs (mycorrhizal penetration sites) partly accounts for the relative absence of Vesicular Arbuscular Mycorrhizae (VAM) in trees with ARD (Mai *et al.*, 1970; Pinochet *et al.*, 1996).

Many researchers have associated ARD with biotic and abiotic factors, acting individually or together. These factors include pathogenic actinomycetes (Westcott *et al.*, 1987; Otto *et al.*, 1993), bacteria (Utkhede *et al.*, 1992; Dullahide

et al., 1994), fungi (Sewell, 1981; Braun, 1995; Mazzola, 1997), dagger nematodes (*Xiphinema americanum* Cobb) (Sultan and Ferris, 1991; Lana *et al.*, 1983), root lesion nematodes [*Pratylenchus penetrans* (Cobb) (Filipjev and Stekhoven) (Hoestra, 1968; Hoestra and Oostenbrink, 1962; Mai *et al.*, 1970), phytotoxins (Benson *et al.*, 1978), nutrient imbalance (Sadowski *et al.*, 1988), and poor soil conditions (Mai and Abawi, 1981). However, the factors vary among apple growing regions and are not fully determined as evidenced by lack of sustainable control measures. Consequently, the list and interactions of agents associated with ARD increase with every new study (Dullahide *et al.*, 1994). Most isolation studies have used soil from single orchards rather than composite soil obtained from more than one orchard and mixed up. Therefore, our immediate objective involved isolating and identifying nematodes and fungi from ARD-infested soils, similar to those previously used to evaluate apple germplasm for resistance or tolerance to ARD. The ultimate objective was to determine pathogens that interacted to cause poor growth of apple seedlings evaluated and in orchards that provided the ARD-infested soils. Separate orchard soils were used for nematodes that

may not survive well the composting process (Merwin and Stiles, 1989). Composted soils were used for fungi under the assumption that fungal reproductive structures can withstand composting. Knowledge of these pathogens might lead to development of sustainable control measures such as plant resistance by screening apple germplasm in ARD-infested composite soil. Resistance or tolerance identified in composite soil rather than in single orchard soils is likely to be broad and durable.

MATERIALS AND METHODS

Apple Hosts

Microshoots of different apple hosts were obtained from the apple rootstock disease-resistance breeding programme at Geneva, New York. Different rootstocks were used to increase chances of isolating different fungi. The tested hosts include T348 'Royal Gala' and T565 'Marshall McIntosh' (MM), each with a chitinase gene, T286MM with a *gus* screenable marker gene, non-transformed MM, non-transformed M.7a, non-transformed 78M26R5 (CG11) and non-transformed *M. spectabilis*-1844. Transgenics and T286MM *gus* were included to determine effects of transgenes, *gus* gene, and transformation process on ARD pathogens. All microshoots were proliferated on modified Murashige and Skoog (1962) basal salts with minimal organics (Sigma Chemical Co., St. Louis, Missouri). Each litre of medium for proliferating non-transformed 78M26R5 (CG11) and non-transformed *M. spectabilis*-1844 was amended with 1 mg each of: thiamine, nicotinic acid, pyridoxine, N⁶-benzyladenosine (BAP), indole-3-butyric acid (IBA), and gibberellic acid (GA₃); 100 mg myo-inositol, 30 g lab grade sucrose, and 7 g Difco Bacto agar. The medium for T565MM Chitinase and T286MM *gus* was amended per litre with 100 mg myo-inositol, 30 g sucrose, 7.5 g agar, 0.3 mg IBA, 1 mg BAP, 0.2 mg GA₃, 0.8 g mg thiamine, 1 mg pyridoxine, 1 mg nicotinic acid, 0.08 mg biotin, 4 mg glycine and 200 µg paromomycin. Non-transformed MM was proliferated on the same medium as for T565MM Chitinase, but without paromomycin. T348RG Chitinase was proliferated on the same medium as for T565MM Chitinase, except IBA and GA₃ were omitted, and 3 mg of kinetin and 0.1 mg 1-naphthalenacetic acid were added. The pH of all media was adjusted to 5.6, the media poured into glass jars or Magenta G7 boxes (Magenta Corp., Chicago, Illinois), and autoclaved for 18 minutes at 121°C and 100 kPa. Culture vessels were sealed with parafilm and placed in a room maintained at 24±2°C with about 70 µmol.m⁻².s⁻¹ of photosynthetically active radiation (PAR) for 16 hours per day. After proliferating for at least four weeks, microshoots were excised and rooted in peat: perlite (1:1, v/v) for about four weeks under mist in a greenhouse maintained at 25°C and 16 hours of light. Rooted plants were grown for 2 weeks with hand-watering before transplanting to ARD-infested soil.

Sources of ARD Soil

Based on previous ARD diagnostic bioassays, orchard surveys, and difficulties encountered in replanting young trees (Pruyne *et al.*, 1994), ARD soil was collected from the

same New York orchards on three occasions. The orchards ranged from old (established) to new (fallowed or recently replanted) sites of the Bartleson, Clarke, Crist, Grainger and Smith farms. Grass was scrapped off the surface of the alleyways in a random pattern; soil scooped with a hand shovel to a 30-cm depth at about 35 sites per orchard, bulked into black polythene bags and stored at 4°C until it was ready for use.

Nematode Isolation and Identification

Nematodes from diverse genera were extracted and identified from the soil. After taking soil samples, separate soils were homogenized (composited) in a motorized mixer and then immediately returned to a cold storage at 4°C. The composited soil was used to grow apple host plants from which fungi were isolated. Nematodes in soil subsamples were extracted for 7 days on modified Bearman pans, and then identified (Zuckerman *et al.*, 1985; Mai and Mullin, 1996).

Isolation and Identification of Fungi

The fungal species targeted were for *Cylindrocarpon*, *Pythium*, *Fusarium*, *Rhizoctonia* and *Phytophthora* previously associated with ARD (Sonderhausen, 1965; Sewell, 1981; Jaffee *et al.*, 1982b; Braun, 1995). About 10 plantlets of each clone were grown in a composite ARD soil, harvested after 10 weeks and processed for fungal isolation. All fungi were isolated and identified by the Plant Pathology Diagnostic Laboratory, except *Phytophthora* species which were isolated and identified by ourselves in the Laboratory at Cornell University, Ithaca, New York. To isolate species of *Cylindrocarpon*, *Pythium*, *Fusarium* and *Rhizoctonia*, roots of apple hosts were washed in running tap water for about two minutes and cut into 1-cm long pieces. Root pieces were surface-disinfected in 10% NaOCl solution for about 2 minutes and rinsed twice in sterilized distilled water. The roots were then blotted dry using a sterile paper, randomly selected and cultured onto selective isolation and identification media. *Pythium* species were isolated by culturing three root pieces from each host root replicate on water agar amended with rifampicin and primaricin (Sigma Chemical Co., St. Louis, Missouri; Jaffee *et al.*, 1982a). Small agaral pieces of unique colonies were subcultured separately within three days on PDA or Corn Meal Agar (CMA) for identification up to genus level and recording isolation frequency (Jaffee *et al.*, 1982a; Reeleder and Brammall, 1994). *Fusarium* and *Cylindrocarpon*, with simpler protocols, were subcultured for identification to species level. Single spores or hyphal tips were subcultured on carnation water agar or PDA for *Fusarium* and PDA for *Cylindrocarpon* (Samuels and Bryford, 1990). *Fusarium* species were identified based on the colour of sporochia (Nelson *et al.*, 1983). *Cylindrocarpon* species were identified based on the presence or absence and size of microconidia, macroconidia and mycelial chlamydospores (Booth, 1966).

Phytophthora species Isolation and Identification

Roots of apple hosts were washed in running tap water for three minutes, surface-disinfected in 70% ethanol for two seconds, rinsed three times in sterilized distilled water,

blotted dry on sterilized paper and cut into 1-cm-long pieces. Nine pieces from each host root replicate were randomly selected and cultured on selective medium containing CMA, primaricin, ampicillin, rifampicin, PCNB and hymexazol (P_5 ARPH) (Mircetich and Mathernon, 1976). After three days at 21°C, emerging colonies were subcultured onto CMA and V-8 juice agar (V8A) for identification and recording isolation frequency (Wilcox and Ellis, 1989). Species were identified based on colony, sporangia and oogonia morphology on CMA and V8A (Wilcox, 1989; Wilcox and Ellis, 1989). Isolation frequency for all fungi was analyzed using the Chi-square test (StatXact, Cytel Software Corp., Cambridge, Massachusetts) to determine whether it varied significantly or not among the different fungi.

RESULTS AND DISCUSSION

Nematodes

Diverse genera of nematodes were present in the ARD soil collected from selected New York orchards (Table 1). The highest population densities were observed for non-parasites, as reported in other studies (Hoestra, 1968; Vrain and Rousselle, 1980; Vrain and Yorston, 1987). Root-lesion nematodes (RLN) and dagger nematodes (DN), both pathogenic to apple were among the nematodes isolated. The frequency of occurrence (10/10, 9/9 and 9/9) for RLN was always 100% (Table 1), whereas (6/10, 6/9 and 4/9) for DN indicated absence of DN in some of the orchards sampled. The presence of RLN in all and of DN in some New York orchards surveyed has been reported by other researchers (Mai and Abawi, 1978; Rosenberger *et al.*, 1983; Driel *et al.*, 1990). Therefore, plant resistance to these two nematodes should be included in strategies designed to combat ARD.

The number of RLN and DN varied by season and orchard (Table 1). In 100 cm³ of separate soil samples, average RLN and DN populations ranged from 56 to 350, and 5 to 58, respectively (Table 1). This variation was consistent with previous reports on seasonal distribution and habits of these nematodes (Constante *et al.*, 1985; Jaffee *et al.*, 1987; Kotcon, 1990; Vrain *et al.*, 1997). Populations within these ranges have frequently been observed in orchards exhibiting ARD (Mai and Abawi, 1978; Vrain and Rousselle, 1980; Merwin and Stiles, 1989; Utkhede *et al.*, 1992). There is no commonly accepted economic injury threshold for RLN and DN in apple. Reported thresholds vary with apple-growing region and other soilborne pathogens acting in association (Hoestra, 1968; Sitepu and Wallace, 1974; Jaffee *et al.*, 1982b; Utkhede and Li, 1988; Dullahide *et al.*, 1994; Utkhede *et al.*, 1992). One estimated threshold ranged from 20 to 50 RLN per 100 cm³ of soil (Hoestra and Oostenbrink, 1962). Hoestra (1968) considered orchards with 50 RLN per 100 cm³ of soil and poor growth of replanted trees as heavily infested. In contrast, injury thresholds for DN can be as low as 1 per 100 cm³ of soil, because of its potential to transmit tomato ringspot virus to apple (Rosenberger *et al.*, 1983; Kotcon, 1990). Therefore, RLN and DN populations in most orchards sampled were relatively high. Based on the substantial stunting that other researchers have reported for

these population densities, we associated RLN, DN and other pathogens in the composite soil with the ARD symptoms observed among apple hosts.

Fungal Species

Fusarium, *Cylindrocarpon*, *Pythium*, *Rhizoctonia*, *Mortierella*, *Trichoderma*, *Penicillium*, *Chaetium*, *Rhizopus* and *Mucor* were isolated from ARD-infested apple roots (Table 2). However, this paper will focus on the pathogenic *Fusarium*, *Cylindrocarpon*, *Pythium* and *Rhizoctonia* previously that have associated with replant disorders (Sewell, 1981; Jaffee *et al.*, 1982a; Mai and Abawi, 1987; Braun, 1995). The frequency of isolation across hosts varied significantly among fungi ($P < 0.05$) (Table 2). *Pythium* species had the highest isolation frequency (29%), followed by four species of *Fusarium* (26%), two *Cylindrocarpon* spp. (12%) and lastly by *Rhizoctonia* sp. (1%) (Table 2). Among the four *Fusarium* species, *F. oxysporum* was the most prevalent (22%) (Table 2). The frequency of *C. lucidum* was higher (7%) than that of *C. destructans* (5%) (Table 2).

Therefore, *Pythium* species may be more aggressive in colonizing apple roots than the other fungi identified. The number of hosts colonized by each fungus varied and was 6 of 6 each for *Pythium* species and *F. oxysporum*, 3 of 6 each for *C. destructans* and *C. lucidum*, and 1 of 6 each for *Pythium* species and *F. acuminatum* and *Rhizoctonia* sp. (Table 2). Thus *Pythium* species and *F. oxysporum* colonized all hosts tested, suggesting that they can cause disease in a wide range of apple rootstocks (Sharma and Gupta, 1989). The eight pathogenic fungi isolated were: *Pythium* species, *F. equiseti*, *F. oxysporum*, *F. solani*, *F. acuminatum*, *C. lucidum*, *C. destructans* and *Rhizoctonia* sp. (Table 2). These fungi varied per host and were 2 of 8 on T348RG Chitinase, 3 of 8 on T565MM Chitinase, 5 of 8 on T286MM *gus*, 4 of 8 on 78M26R5 (CG11), 6 of 8 on 'Marshall McIntosh', and 3 of 8 on *M. spectabilis*-1844 (Table 2). Similar variation in fungal infection has been observed among apple rootstocks (Jones and Aldwinckle, 1990). It appears diverse hosts and the composite soil enabled us to isolate different fungi. Both transgenic and non-transgenic apple hosts were infested by several fungi. Endochitinase, the *gus* gene, and gene-transfer process did not prevent fungal infection. Several *Fusarium* species, including *F. equiseti*, *F. oxysporum* and *F. solani* isolated have been associated with ARD in the United States (Utkhede and Smith, 1994). *F. acuminatum*, which has not been associated with ARD before was also isolated. However, *Fusarium* species cause considerable damage in nursery plants (Mazzolla, 1997). *Cylindrocarpon* species previously associated with ARD include *C. lucidum*, *C. scoparium*, *C. radicola*, *C. heteronema* and *C. destructans* (Jaffee *et al.*, 1982a; Braun 1995). *Cylindrocarpon lucidum* caused lower biomass, discolouration, black lesions and extensive cortical decay symptoms on roots; these symptoms were more severe when *P. irregulare* interacted with *C. lucidum*.

TABLE 1: Population of nematodes extracted from soils for sampled New York orchards

Year 1 Season 1 (Spring)	Frequency	Orchard										Average ± SD
		1 ^x	2 ^x	3 ^x	4	5	6	7	8	9 ^x	10	
Non-parasites	10/10	478	601	1443	1794	960	624	476	280	306	266	723±521
<i>Pratylenchus penetrans</i>	10/10	40	39	51	78	72	72	68	84	73	38	62±18
<i>Heterodera</i> sp.	8/10	52	0	0	26	72	120	34	28	18	152	50±51
<i>Pratylenchus</i> sp.	8/10	82	26	45	26	192	0	34	0	124	76	61±60
<i>Cricemella</i> sp.	7/10	0	173	102	26	24	0	0	280	180	76	86±97
<i>Filenchus</i> sp.	7/10	54	26	15	0	120	48	0	84	18	0	37±40
Predators	6/10	14	108	17	0	144	24	0	0	11	0	32±51
<i>Xiphinema americanum</i>	6/10	40	0	17	0	192	0	0	224	65	38	58±83
<i>Costenichus</i> sp.	4/10	232	18	32	0	0	0	0	0	54	0	34±72
<i>Aphelenchus</i> sp.	4/10	0	18	17	0	24	48	0	0	0	0	11±16
<i>P. silenichus</i> sp.	3/10	176	0	60	78	0	0	0	0	0	0	31±59
<i>Helicotylenchus</i> sp.	2/10	0	0	0	15	0	0	0	0	0	0	4±10
<i>Tylenchus</i> sp.	4/10	0	0	0	0	24	0	34	140	126	0	32±54
<i>Tylenchorhynchus</i> sp.	2/10	0	0	0	26	0	0	0	56	0	0	8±19
Year 1 Season 2 (Fall)												
<i>Pratylenchus penetrans</i>	9/9	115	254	132	318	352	120	635	550	678		350±222
<i>Xiphinema americanum</i>	9/9	20	21	0	49	64	0	18	0	7		20±23
Year 2 Season 2 (Fall)												
<i>Pratylenchus penetrans</i>	9/9	1 ^v	2 ^v	3	4 ^w	5 ^v	6 ^v	7 ^v	8 ^w	9 ^v		56±36
<i>Xiphinema americanum</i>	4/9	61	24	39	71	77	49	10	134	36		5±7

^{z1} = Bartleson I, 2 = Bartleson II, 3 = Clarke, 4 = Crist I, 5 = Crist II, 6 = Grainger I, 7 = Grainger II, 8 = Smith II, 9 = Smith I, 10 = Trapani. I = fallowed or recently planted. II = Established. SD = Standard Deviation

^y Extracted by sucrose floatation, the rest by modified Bearman pie pans (Zuckerman et al., 1985)

^{x,w,v} Average of two, three and four samples, respectively

TABLE 2: Isolation frequency (percentage ±standard deviation) of fungi infesting roots of various apple hosts grown in a composite ARD-infested soil

Fungi	Apple hosts										% Frequency/ fungus	Hosts infested
	T348RG Chitinase	T565MM Chitinase	T286MM gus	78M26R5 (CG11)	Marshall McIntosh	<i>M. spectabilis</i> 1844						
<i>Pythium</i> species ^z	43±40 ^y	31±39	33±15	46±31	6±10	13±22	29					6 of 6
<i>Fusarium oxysporum</i> ^z	32±16	6±10	40±11	5±8	26±23	25±8	22					6 of 6
<i>Mortierella</i> sp.	17±29	17±17	0±0	21±18	22±9	25±8	17					5 of 6
<i>Cylindrocarpum lucidum</i> ^z	0±0	0±0	5±8	15±15	6±10	22±25	7					3 of 6
<i>Trichoderma</i> sp.	0±0	11±19	5±8	0±0	11±10	6±10	6					4 of 6
<i>C. destructans</i> ^z	0±0	14±13	0±0	0±0	9±9	0±0	5					3 of 6
Unknown	8±14	6±10	0±0	0±0	9±9	0±0	4					3 of 6
<i>Penicillium</i> sp.	0±0	6±10	11±10	0±0	6±10	6±10	3					3 of 6
<i>F. equiseti</i> ^z	0±0	0±0	0±0	0±0	0±0	0±0	2					1 of 6
<i>Chaetomium</i> sp.	0±0	11±19	0±0	0±0	0±0	0±0	2					1 of 6
<i>Fusarium solani</i> ^z	0±0	0±0	7±12	5±8	0±0	0±0	1					1 of 6
<i>Fusarium acuminatum</i> ^z	0±0	0±0	0±0	0±0	0±0	0±0	1					1 of 6
<i>Rhizopus</i> sp.	0±0	0±0	0±0	8±14	0±0	0±0	1					1 of 6
<i>Rhizoctonia</i> sp. ^z	0±0	0±0	0±0	0±0	6±10	0±0	1					1 of 6
<i>Mucor</i> sp.	0±0	0±0	0±0	0±0	0±0	0±0	1					1 of 6
Pathogenic fungi	2 of 8	3 of 8	5 of 8	4 of 8	6 of 8	3 of 8	* ^w					1 of 6

^zIndicates fungi pathogenic to apple

^yEach frequency is a mean of three replicate root systems. Total frequency is 100 for each host

^xPercentage isolation frequency for each fungus is average of six hosts

^wIndicates significant difference in percentage isolation frequency among fungi (Chi-square test, P<0.05)

TABLE 3: Isolation frequency (percentage ± standard deviation) for *Phytophthora* species infesting roots of various apple hosts in a composite ARD-infested soil

Fungi	Apple hosts						M. spectabilis- 1844	% Frequency/ fungus ^x	Hosts infested
	T348RG Chitinase	T565MM Chitinase	M.7a (Control)	78M26R5 (CG11)	'Marshall McIntosh'				
None	67±58 ^y	66±34	55±51	33±0	33±33	22±19	46	6 of 6	
Other fungi	11±19	33±33	33±58	22±19	55±39	33±0	31	6 of 6	
<i>Phytophthora cryptogea</i> ^z	5±9	0±0	0±0	33±33	0±0	22±38	10	3 of 6	
<i>Phytophthora cactorum</i> ^z	0±0	0±0	0±0	0±0	11±19	11±19	4	2 of 6	
<i>Phytophthora cambivora</i> ^z	11±19	0±0	0±0	11±19	0±0	11±19	4	2 of 6	
<i>Phytophthora megasperma</i> ^z	5±9	0±0	0±0	0±0	0±19	11±19	3	2 of 6	
<i>Phytophthora</i> sp. ^z	0±0	0±0	11±19	0±0	0±0	0±0	2	1 of 6	
<i>Phytophthora</i> spp.	3 of 5	0 of 5	1 of 5	2 of 5	1 of 5	3 of 5	* ^w		

^zIndicates *Phytophthora* species pathogenic to apple

^yEach frequency is a mean of three replicate root systems. Total frequency is 100 for each host

^xPercent isolation frequency for each *Phytophthora* species is average of six hosts

^wIndicates a significant difference in percentage isolation frequency among species (Chi-square test P <0.05)

Pythium species that have been isolated from orange-coloured lesions on apple roots previously grown in ARD soil varied with the geographic region and included *Pythium irregular*, *Pythium sylvaticum*, *Pythium ultimum*, *Pythium intermedium* and others (Sewell, 1981; Dullahide et al., 1994; Braun, 1995). It is possible that the *Pythium* isolates included some of these species. *Pythium ultimum* has been associated with collar rot of apple (Sharma and Gupta, 1989).

A few *Rhizoctonia* species have been isolated from ARD soils and roots. Mazzolla (1997) reported that *R. solani* isolates in anastomosis groups 5 and 6 were the most dominant and pathogenic in Washington State soils and they caused severe symptoms of ARD in artificial inoculations. Since *Rhizoctonia* sp. was isolated from only one root system of 'Marshall McIntosh', it may be less prevalent in New York orchards. These results agree with observations that *Rhizoctonia* is primarily a damping-off and root rot fungus in seed beds and nurseries (Mazzolla, 1997).

Phytophthora species

Phytophthora cryptogea, *Phytophthora cactorum*, *Phytophthora cambivora*, *Phytophthora megasperma* and *Phytophthora* sp., were isolated from apple hosts tested (Table 3). The isolation frequency of *Phytophthora cryptogea* was highest (10%) followed by *Phytophthora cambivora* and *Phytophthora cactorum* (4% each), *Phytophthora megasperma* (3%) and lastly *Phytophthora* sp. (2%) (Table 3).

Phytophthora cryptogea was also isolated from 3 of 6 hosts, while each of the other species was found on 2 of 6 hosts. *Phytophthora* sp. occurred on 1 of 6 hosts (Table 3). Although *Phytophthora cryptogea* was the most prevalent, all the *Phytophthora* species isolated have previously been isolated from ARD-infested roots (Sitepu and Wallace, 1974; Utkhede et al., 1992). They also cause crown rot in apple (Jeffers et al., 1982; Jones and Aldwinckle, 1990).

The number of species per host was zero on T565MM Chitinase, 1 of 5 each on M.7a and 'Marshall McIntosh' controls, 2 of 5 on 78M26R5 (CG11), and 3 of 5 on T348RG Chitinase and *M. sectabilis*-1884 (Table 3). Similar variation has been observed among apple rootstocks, where M.9 was the most resistant, M.7 and M.7a were moderately susceptible and M.26 was moderate to very susceptible to *Phytophthora* (Jones and Aldwinckle, 1990).

CONCLUSION

The ARD soil from New York orchards had both fungi and nematodes, some of which are known to be pathogenic to apple. These pathogens and others not isolated most likely interacted to cause ARD symptoms in the apple hosts. Therefore, composite soil from ARD-infested apple orchards are a good source of inoculum for screening *Malus* germplasm for resistance or tolerance to mixed ARD pathogens. Resistance to a combination of pathogens may be more durable than resistance to single pathogens.

RECOMMENDATIONS

Whenever unthrifty growth is observed in soil especially that under perennial fruits and continuously cultivated

annual crops, similar diagnostic isolation of multiple pathogens should be performed. In developing management strategies, germplasm for the crop in question should be evaluated in multiple sites or in composite soil in order to identify germplasm with broad-spectrum resistance or tolerance to the pathogens. Another technique than can accelerate development of broad-spectrum resistance is gene transfer.

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