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FUSARIUM ROOT DISEASE OF BARE ROOT 1-0 WESTERN WHITE PINE AND DOUGLAS-FIR SEEDLINGS USDA FOREST SERVICE NURSERY, COEUR D'ALENE, IDAHO

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ABSTRACT

Isolations from roots of scattered diseased 1-0 western white pine and Douglas-fir bare root seedlings at the USDA Forest Service Nursery, Coeur d'Alene, Idaho yielded high levels of *Fusarium*, particularly *F. oxysporum*. Fairly high levels of *F. sporotrichioides* and *F. acuminatum* were also isolated from roots of diseased white pine seedlings. Since all seedlings were grown in soil previously fumigated with dazomet, it is likely that much pathogen inoculum occurred on sown seeds, particular in white pine beds where resident levels of soil *Fusarium* populations were low. It is important to monitor commonly-used seedlots for seed-borne pathogen potential and subject highly contaminated lots to surface sterilization treatments.

INTRODUCTION

Root diseases caused by *Fusarium* spp. are probably the most important production-limiting diseases at the USDA Forest Service Nursery in Coeur d'Alene, Idaho. These diseases are especially important during the first growing season on bare root conifer seedlings (James 2001, 2002). Seedborne *Fusarium* is often associated with damping-off losses during seed germination and seedling establishment (James 1986, 1987) and soil-borne inoculum is important in causing both damping-off and root diseases of older seedlings (James et al. 1991). Generally, seedlings surviving the first growing season are usually not adversely affected during their second growing season. Although all conifer species may be affected by *Fusarium* diseases, Douglasfir (*Pseudotsuga menziesii* Franco var. *glauca* [Mayr.] Sudw.) and western white pine (*Pinus monticola* Dougl.) are quite susceptible and often severely impacted (James et al. 2004a, 2004b).

Near the end of the first growing season, stands of bare root western white pine seedlings were extremely sparse (figure with recently-killed seedlings 1). interspersed randomly throughout stands (figure 2). Intermixed within seedling stands were extensive weed populations (figure 3). At the same time, in an adjacent field, extensive mortality of 1-0 Douglas-fir seedlings was evident (figure 4). Recently-killed seedlings were interspersed with healthyappearing seedlings.

Since similar disease had previously been reported at the Coeur d'Alene Nursery (James 1993, 2000; James et al. 2004a, 2004b), it was suspected that root pathogens were probably responsible for mortality and likely contributed to the sparse seedling stands. An evaluation was conducted to confirm identity of associated organisms and determine soil populations of potential pathogens within affected areas.

MATERIALS AND METHODS

Three sets of isolations were made: from diseased and healthy-appearing seedlings, from soil surrounding the roots of diseased seedlings, and from soil obtained from the rhizosphere of diseased seedlings. All isolations were made onto a selective agar medium for Fusarium and closely-related fungi (Komada 1975). Isolations were made from roots of seven and eleven recentlykilled Douglas-fir and western white pine seedlings, respectively. Similar isolations were made from two and three healthy-appearing Douglas-fir and western white pine seedlings, respectively. Selected seedlings were carefully extracted from soil to include as much of their root systems as possible. Seedling roots were carefully washed under running tap water and dissected into pieces approximately 5 mm in length; the entire attached root system of each seedling was sampled (10-21)root pieces sampled per seedling). All root pieces were surface sterilized in an aqueous solution of 0.525% sodium hypochlorite (10% common household bleach), rinsed in sterile distilled water, blotted dry, and placed on the selective agar medium. Fifteen root pieces were placed on agar within each petri plate. Roots were incubated at about 24°C under diurnal cycles of cool, fluorescent light for 7 days. Associated fungi were identified to genus based on spore morphology (Barnett and Hunter 1998).

Selected colonies of suspected pathogens were transferred to carnation leaf (Fisher et al. 1982) and potato dextrose agar (PDA) for identification. *Fusarium* species were identified using the taxonomy of Nelson et al. (1983). The percent of sampled seedlings infected and the extent of root system colonized by specific potential pathogens were calculated.



Figure 1. Sparse stands of 1-0 bare root western white pine seedlings - USDA Forest Service Coeur d'Alene Nursery, Idaho.



Figure 2. Mortality of 1-0 bare root western white pine seedlings - USDA Forest Service Nursery, Coeur d'Alene, Idaho. Affected seedlings were randomly scattered throughout seedbeds.



Figure 3. Recently-killed 1-0 bare root western white pine seedling surrounding by high populations of groundcover weeds - USDA Forest Service Nursery, Coeur d'Alene, Idaho.



Figure 4. Several recently-killed 1-0 bare root Douglas-fir seedlings - USDA Forest Service Nursery, Coeur d'Alene, Idaho.

Soil samples were collected adjacent to diseased seedlings; three and four samples were obtained in Douglas-fir and western white pine beds, respectively. Samples were collected to a depth of about 20 cm using a standard soil probe (about 5 mm in diameter).

Each soil sample consisted of cores from four probe sub-samples (collected at the four cardinal directions around a diseased seedling); the sub-samples were collated and mixed together to form a larger sample. Soil was placed into plastic bags, kept refrigerated, and transported to the laboratory for analysis.

dilutions (Hildebrand Standard and Dinkel 1988: James and Beall 1999: James et al. 1990, 1996) were conducted on the soil samples. Soil from each sample was initially sieved (2 mm sieve) to remove rocks, pieces of organic matter, and soil aggregates. From each sample, an approximate 5 g subsample was oven-dried at about 100°C for at least 24 hours until sample weight had stabilized. Oven-dry weight was then calculated to provide a standard for sample comparisons. For assays of Fusarium and Trichoderma populations, 0.05 g of field-moist soil was combined with 10 ml of 0.3 percent water agar and thoroughly mixed. One ml of solution was placed on each of three plates of the selective agar medium and spread uniformly. Plates were incubated as described above. Fusarium and Trichoderma colonies were identified by their morphology on the selective medium; populations were calculated as number of colony-forming units (cfu) per g of oven-dried soil. Selected Fusarium isolates were identified as described above.

Soil adhering to the roots of diseased seedlings (rhizosphere soil) was also assayed. Roots from three each of Douglas-fir and western white pine seedlings displaying above-ground disease symptoms (necrotic foliage) were immersed in 100 ml of sterile, distilled water and thoroughly agitated to remove rhizosphere soil. The solution was thoroughly blended and 1 ml of the solution was placed on three plates of selective agar medium (for each sample) and spread uniformly. Plates were incubated as described above and the average number of Fusarium colonies per ml of rhizosphere soil solution was calculated.

RESULTS AND DISCUSSION

All sampled diseased western white pine and Douglas-fir seedlings were infected with *Fusarium* spp. (table 1). Most healthy-appearing seedlings were likewise infected. Root systems of diseased seedlings were extensively colonized by these potential pathogens. However, root systems of healthy white pine were much less colonized by *Fusarium* spp. compared to those of healthy Douglas-fir.

Soil surrounding roots of diseased Douglas-fir seedlings contained fairly high levels of *Fusarium* propagules (table 2), whereas those near white pine seedlings had much lower populations. Higher *Fusarium* levels generally corresponded to lower *Trichoderma* populations, resulting in lower ratios of *Trichoderma* to *Fusarium* populations. Lower ratios may denote decreased potential for disease suppressiveness by species of *Trichoderma* which can potentially exert biological control (James and Beall 1999; Papavizas 1985).

Seven different Fusarium species were isolated from either roots of white pine and Douglas-fir seedlings or adjacent soil (table 3). By far the most commonly-associated species was F. oxysporum Schlecht.. Three different morphotypes of this large, geneticallydiverse species were encountered. Morphotypes were separated on the basis of colony morphology on PDA, i.e., extent of aerial or suppressed mycelium (pionnotal), pigmentation, and sporodochial production. Isolates of Fusarium oxysporum may exhibit a wide range of pathogenic potential on conifer seedlings (Gordon and Martyn 1997; James et al. 1991; Nelson et al. 1983).

Because pathogenic and non-pathogenic strains appear similar morphologically (Bao et al. 2002; Gordon and Martyn 1997), it is unknown what proportion of the isolates obtained in this investigation were pathogens. It is suspected that most from roots of isolates seedlings exhibiting disease symptoms were pathogens, although pathogenic strains may occur on roots of seedlings without above-ground disease symptoms (James et al. 2000). Recent genetic work (Stewart et al. 2004) indicated that pathogenic and non-pathogenic F. oxysporum strains from forest nurseries may be separated on the basis of molecular genetic markers. If further work confirms this, rapid analysis of pathogenic populations within nurseries may be possible in the future.

Table 1. *Fusarium* infection and colonization of roots from diseased and healthyappearing 1-0 bare root western white pine and Douglas-fir seedlings - USDA Forest Service Nursery, Coeur d'Alene, Idaho.

Sample Category	Western White Pine		Douglas-fir		
	Diseased	Healthy	Diseased	Healthy	
Number of Seedlings Sampled	11	3	7	2	
Percent of Seedlings Infected ¹	100	67	100	100	
Number of Root Pieces Sampled	155	55	105	37	
Percent of Root Pieces Colonized ¹	87.1	16.4	97.1	100	

Infected and colonized by Fusarium spp.

Table 2. Populations of *Fusarium* and *Trichoderma* spp. within nursery soil adjacent to diseased western white pine and Douglas-fir seedlings - USDA Forest Service Nursery, Coeur d'Alene, Idaho.

Fungal Genera	Soil Populations [cfu/g]			
	Western White Pine	Douglas-fir		
Fusarium	154	1519		
Trichoderma	291	91		
T/F Ratio ¹	1.89	0.06		

¹ Ratio of *Trichoderma* to *Fusarium* populations; higher numbers denote greater potential for disease suppression by *Trichoderma* spp.

Table 3. *Fusarium* species colonizing 1-0 bare root western white pine and Douglas-fir seedlings and adjacent soil - USDA Forest Service Nursery, Coeur d'Alene, Idaho.

Fusarium Species ¹	Western White Pine ²		Douglas-fir ²		Soil ³
	Diseased	Healthy	Diseased	Healthy	1. 193
FOXY-1	36.8	0	80.9	89.2	67.1
FOXY-2	5.2	0	30.5	54.0	10.5
FOXY-3	3.9	0	11.4	10.8	6.6
ALL FOXY	45.8	0	85.4	100.0	84.2
FAVE	0	0	9.5	0	11.8
FACU	23.9	5.4	5.7	0	4.0
FSPO	27.7	10.9	0	0	0
FCUL	0	0	5.7	0	0
FEQU	0.6	0	0	0	0
FSAM	0	7.3	0	0	0

¹ FOXY = Fusarium oxysporum [numbers denote different morphotypes]; FAVE = F. avenaceum; FACU = F. acuminatum; FSPO = F. sporotrichioides; FCUL = F. culmorum; FEQU = F. equiseti; FSAM = F. sambucinum.

² Percent of sampled root pieces colonized by particular Fusarium species.

³ Percent of *Fusarium* isolates obtained from all soil samples.

All the *Fusarium* species encountered have previously been isolated at the Coeur d'Alene Nursery (James et al. 1989). They are commonly found within nursery soil (James 2000b, 2004; James and Perez 1999; James et al. 1996, 2004a, 2004b) as well as on conifer seedling roots (James and Perez 1999; James et al. 2004a, 2004b). Fusarium sporotrichioides Sherb. and F. acuminatum Ell. & Ev. were isolated at relatively high levels from the roots of diseased white pine seedlings (table 3). These species probably are saprophytes and may not be capable of eliciting seedling disease unless hosts are extensively stressed (James 2000a; James and Perez 1999). However, it is possible that one or more of these species may affect disease development by influencing the ability of pathogenic strains of F. oxysporum to cause disease (Hoff et al. 2004; James et al. 1991). Some Fusarium strains are rootcolonizing endophytes of conifer seedlings and may be present without adversely affecting host physiology (Bloomberg 1966; Hoff et al. 2004). They would be routinely isolated from surface-sterilized roots (Hoff et al. 2004; Stone et al. 2000), but not necessarily act as pathogens. Further work is required to elucidate their potential roles in seedling disease etiology.

Soil within which diseased seedlings were found had been fumigated with dazomet (Basamid®) prior to sowing. Therefore, most potentially-pathogenic fungi had probably been killed by fumigation. Pathogens isolated from diseased seedling roots were most likely introduced on contaminated seed.

Most pathogens on conifer seed are located externally on seedcoats as passive contaminants (James 1986. 1987). If they can be satisfactorily removed, disease potential to young seedlings is greatly decreased (James 1986). Pre- or post-stratification running water rinses for a minimum of 48 hours greatly reduce seed can surface contamination (Campbell and Landis 1990; Dumroese et al. 1988; James 1986, 1987), particularly if seeds are agitated during rinsing. However, highly-contaminated seeds may need treatment with surface sterilants to adequately remove most pathogen propagules (Campbell and Landis 1990; James 1986, 1987). Chemical treatments

are often operationally used on conifer seeds; the two most effective are sodium hypochlorite (bleach) and hydrogen peroxide. Bleach treatment prior to running water rinses has become standard for seeds of many conifer species at some nurseries (Wenny and Dumroese 1987). Hydrogen peroxide is used less frequently, but can be effective as long as chemical concentrations are insufficient to elicit phytotoxicity on young seedling germinants (Barnett 1976; Carter and Jones 1962). Some growers have treated seeds with fungicides (Berbee et al. 1953; Bloomberg and Trelawny 1970). Although results have varied, most fungicides are usually not recommended for seed treatment because of their high potential to reduce germination and cause damage to young, succulent germinants (Cayford and Waldon 1967; Fraser and Adams 1980).

In most cases, conifer seedlots are not routinely screened in the United States to determine level of contamination with potential pathogens. Such tests are becoming common elsewhere, particularly in Canada and parts of Europe. Testing widely-used seedlots for presence of pathogens will help growers select the best lots for sowing as well as concentrate their treatment efforts on those lots exhibiting the greatest contamination (Campbell and Landis 1990). Reducing seedborne pathogens will greatly improve seed germination, seedling establishment, and performance in both bare root and container nurseries.

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