

Short communication

## ***Fusarium redolens* f.sp. *asparagi*, causal agent of asparagus root rot, crown rot and spear rot**

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### **Abstract**

Two *Fusarium* species, *F. oxysporum* f.sp. *asparagi* and *F. proliferatum*, are known to be involved in the root and crown rot complex of asparagus. We have investigated reports on the involvement of *F. redolens*, a third species, which until recently was considered conspecific with *F. oxysporum* because of morphological similarities. RFLP analysis of the rDNA internal transcribed spacer region and AFLP fingerprinting identified eight strains from asparagus unambiguously as *F. redolens*. Four of these were tested and found to be pathogenic to asparagus either in this study (two strains) or in a previous one in which they were classified as *F. oxysporum* (three strains). Disease symptoms and disease development were the same as with *F. oxysporum* f.sp. *asparagi* and *F. proliferatum*. Present data and literature reports identify *F. redolens* as a host-specific pathogen involved in root, crown and spear rot of asparagus. The pathogen is formally classified as *F. redolens* Wollenw. f.sp. *asparagi* Baayen.

Several pathogenic *Fusarium* species are associated with asparagus diseases. Foot and stem rot is caused by *F. culmorum* (W.G. Smith) Sacc., while root and crown rot are generally ascribed to *F. oxysporum* Schlecht.:Fr. f.sp. *asparagi* Cohen & Heald and *F. proliferatum* (Matsushima) Nirenb. (Elmer, 1990, 1991, 1995; Elmer and Stephens, 1989; Blok and Bollen, 1995, 1997; Elena and Kranias, 1996). Spear rot of asparagus has been attributed to *F. redolens* Wollenw., a species that has also been reported to be associated with root and crown rot of asparagus (Gerlach, 1961; UnterEcker, 1972; Cheah, 1986; Falloon and Tate, 1986; Gordon-Lennox and Gindrat, 1987; Sadowski and Knaflewski, 1990).

*Fusarium redolens* was described by Wollenweber (1913) and maintained in his following publications (Wollenweber, 1916–1935, 1931; Wollenweber

and Reinking, 1935). Outside Germany, the country where Wollenweber's taxonomic system was followed (Gerlach and Nirenberg, 1982), *F. redolens* has long been considered conspecific with *F. oxysporum* (Snyder and Hansen, 1940; Nelson et al., 1983) or at best a variety of this species, *F. oxysporum* Schlecht.:Fr. var. *redolens* (Wollenw.) Gordon (Gordon, 1952; Booth, 1971, 1975). Morphological distinction of *F. redolens* from *F. oxysporum* is indeed problematic due to the presence of intermediate forms (Baayen and Gams, 1988). Unambiguous distinction between both species has only recently become possible through molecular means such as RFLP analysis of the rDNA internal transcribed spacer (ITS) region and gene sequence analysis (Waalwijk et al., 1996a,b; O'Donnell et al., 1998). As a consequence, it is uncertain whether former reports on the involvement

of *F. redolens* in asparagus rot are reliable, and also, whether published reports on *F. oxysporum* f.sp. *asparagi* pertain to this taxon, or to *F. redolens*, or to members of both species.

The involvement of *F. redolens* in the spear, root and crown rot complex of asparagus was evaluated in the present study using ITS-RFLP and AFLP analyses as well as pathogenicity tests.

### Identification of strains as *Fusarium redolens*

Isolates from asparagus received as *F. oxysporum* and/or *F. redolens* were obtained from The Netherlands, Germany, Poland, Italy, Greece, U.S.A., South Africa, and New Zealand. Among these were members of 17 vegetative compatibility groups (VCG 1–9 and VCG 11–18) of *F. oxysporum* f.sp. *asparagi* present in The Netherlands (Blok and Bollen, 1997), and members of *F. oxysporum* f.sp. *asparagi* VCG 1001–1008 present in the U.S.A. (Elmer and Stephens, 1989). In total 126 strains were examined. Additionally, three strains of *F. proliferatum* were selected as a reference.

Fungal strains were grown in potato dextrose broth (Difco) at 25 °C for 3 days. DNA was isolated

from lyophilized mycelium as reported previously (Waalwijk et al., 1996a). The ITS region was amplified with primers ITS-1 and ITS-4, and amplification products were digested individually with *Hinf*I, *Alu*I, *Mun*I, and *Rsa*I and analyzed on 1.5% agarose gels as reported previously (Waalwijk et al., 1996a). Isolates were identified by their characteristic ITS restriction fragment length profile (Table 1). ITS-RFLP profiles characteristic of *F. redolens* were obtained for eight strains from The Netherlands, Germany and Poland (Table 2); 117 of the remaining 118 strains were classified as *F. oxysporum* (not shown). Strain IPO 97-10 from New Zealand, received as *F. redolens* under the code 4869-95, proved to belong to *F. proliferatum* and indeed

Table 1. Restriction fragment length sizes of amplified DNA from the ITS region of *F. oxysporum*, *F. redolens* and *F. proliferatum* restricted with *Hinf*I, *Alu*I, *Mun*I, and *Rsa*I

	<i>Hinf</i> I	<i>Alu</i> I	<i>Mun</i> I	<i>Rsa</i> I
<i>F. oxysporum</i>	90, 180, 265	60, 135, 350	545	545
<i>F. redolens</i>	265, 290	560	560	560
<i>F. proliferatum</i>	265, 290	140, 420	70, 490	100, 460

Table 2. Strains of *F. redolens*, *F. oxysporum* and *F. proliferatum* from asparagus plants or from asparagus production field soil. Strains found to belong to *F. oxysporum* and exclusively tested by ITS-RFLP are not listed

Isolate	Origin	Source	ITS-RFLP*	AFLP*	Bioassay*
<i>F. redolens</i>					
CWB1	The Netherlands	W.J. Blok	+	+	+
CWB33	The Netherlands	W.J. Blok	+	+	–
CWB42	The Netherlands	W.J. Blok	+	+	–
IPO 97-13	The Netherlands	J.T.K. Poll	+	+	+
IPO 97-143	The Netherlands	R.P. Baayen	+	–	–
IPO 97-158	The Netherlands	R.P. Baayen	+	+	–
DSM 62380	Germany	W. Gerlach	+	+	–
KFL63	Poznan, Poland	H. Kwasna	+	–	–
<i>F. oxysporum</i>					
MA25 (= FGSC 6607)	U.S.A.	W.H. Elmer	+	+	–
IPO 97-12	The Netherlands	J.T.K. Poll	+	+	–
IPO 97-14	The Netherlands	J.T.K. Poll	+	–	+
IPO 97-102	The Netherlands	R.P. Baayen	+	–	+
IPO 97-147	The Netherlands	R.P. Baayen	+	+	–
IPO 97-172	The Netherlands	R.P. Baayen	+	+	–
<i>F. proliferatum</i>					
IPO 97-03	The Netherlands	J.T.K. Poll	+	+	–
IPO 97-10	New Zealand	E.H.C. McKenzie	+	+	+
M6371	U.S.A.	W.H. Elmer	+	+	–
IPO 97-101	The Netherlands	R.P. Baayen	+	+	+

\*Included (+) or not included (–) in the analyses.

was found to produce the characteristic microconidial chains of this species. Three strains presently identified as *F. redolens* (CWB1, CWB33 and CWB42) had previously been considered to belong to *F. oxysporum* f.sp. *asparagi* (Blok and Bollen, 1997) and represent the three main vegetative compatibility groups (VCG 1, VCG 9 and VCG 12) found in the latter study among 24 isolates from asparagus in The Netherlands.

Six isolates of *F. redolens* identified as such by ITS-RFLP were further characterized by AFLP fingerprinting, along with four isolates each of *F. oxysporum* and *F. proliferatum*. AFLP analyses were performed as described previously (Baayen et al., 2000). Briefly, DNA was digested with *Eco*RI and *Msp*I, followed by ligation of adapters and non-selective amplification with zero primers *Eco*00 (5'GACTGCGTACCAATTC) and *Msp*00 (5'GATGAGTCCTGATCGG). Selective PCR was performed on the amplification product with Cy5-labelled fluorescent *Eco*20 primer (5'GACTGCGTACCAATTCGC) and either *Msp*15 (5'GATGAGTCCTGATCGGCA) or *Msp*16 (5'GATGAGTCCTGATCGGCC) primer. Denatured product was loaded on 6% polyacrylamide gels and run on the automatic sequencer ALFexpress (Pharmacia) at 1500 V, 60 mA, 25 W and 55 °C. A fluorescently labelled 50 bp ladder (Pharmacia) was used as a standard. AFLP patterns were analyzed with ImageMaster software (Amersham Pharmacia Biotech). Only reproducible bands were incorporated in the analysis. A similarity matrix was constructed using the method of Nei and Li (1979). UPGMA cluster analysis of binary data was performed with Treecon software (Van de

Peer and De Wachter, 1994). UPGMA similarity analysis supported the distinction of *F. redolens* as a species distinct from *F. oxysporum* and *F. proliferatum* with 100% bootstrap support (Figure 1).

### Pathogenicity of *Fusarium redolens* to asparagus

The pathogenicity of CWB1, formerly considered to belong to *F. oxysporum* f.sp. *asparagi* but presently shown to belong to *F. redolens*, is well known (Blok and Bollen, 1997). CWB1 is pathogenic to asparagus, but not to 20 other plant species including *Allium* spp., *Anethum graveolens*, *Apium graveolens*, *Beta vulgaris*, *Brassica oleracea*, *Chenopodium album*, *Daucus carota*, *Linum usitatissimum*, *Lolium perenne*, *Medicago sativa*, *Phaseolus vulgaris*, *Poa annua*, *Stellaria media*, *Vicia faba*, and *Zea mays*. Occasionally, pea and lupin may show mild symptoms of foot rot and root rot with CWB1. The same pattern of host specificity was observed for strain CWB 7 (Blok and Bollen, 1997) of which the classification as *F. oxysporum* f.sp. *asparagi* was presently confirmed by ITS-RFLP. The above-mentioned reports by Blok and Bollen (1997) on the host specificity of *F. redolens* CWB1 are concordant with earlier reports by Graham (1955) for *F. redolens* from asparagus. Both *F. oxysporum* f.sp. *asparagi* and *F. redolens* from asparagus thus appear to be restricted in pathogenicity to asparagus.

Apart from its possible involvement in the asparagus rot complex, *F. redolens* has been reported to be responsible for rot and wilt diseases in, among

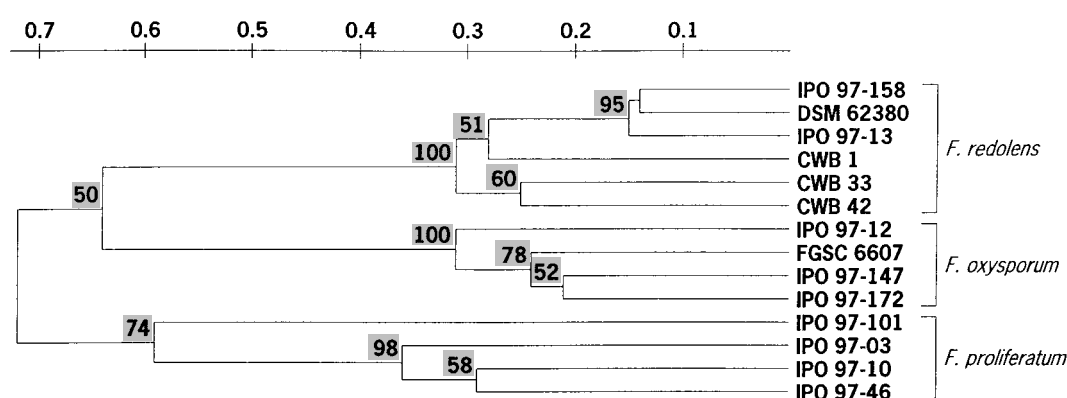


Figure 1. UPGMA similarity analysis of six strains of *F. redolens* from asparagus, and four strains each of *F. oxysporum* f.sp. *asparagi* and *F. proliferatum*. Dissimilarity percentages between isolates are given on the X-axis; bootstrap support percentages (1000 replicates) are indicated at the nodes.

others, carnation, conifers, flax, pea, and spinach (Gerlach and Pag, 1961). In several of these diseases the *F. redolens* strains involved are also host specific. Two formae speciales have been formally published, *F. redolens* f.sp. *dianthi* (Gerlach and Pag, 1961; Baayen et al., 1997, 1999) and *F. redolens* f.sp. *spinaciae* (Sherb.) Subramanian (Subramanian, 1971). Isolates of *F. redolens* f.sp. *dianthi* are not pathogenic to asparagus, and exclusively infect *Dianthus* species (Gerlach and Pag, 1961). The pattern of host specificity of *F. redolens* f.sp. *spinaciae* is not well documented in the original description (Hungerford, 1923).

Pathogenicity of *F. redolens* CWB1 to asparagus was confirmed in the present study for asparagus seedlings (cv. Gijnlim) grown in a growth chamber on water agar in culture tubes at 23 °C and 16 h illumination and inoculated 14 days after germination with a 1 × 1 mm agar piece from a Petri dish culture per seedling as described previously (Blok and Bollen, 1997). After 30 days, CWB1 was equally pathogenic to asparagus as *F. oxysporum* f.sp. *asparagi* IPO 97-102 and *F. proliferatum* IPO 97-101 (Figure 2a), with identical root infection symptoms. Noninoculated control plants remained

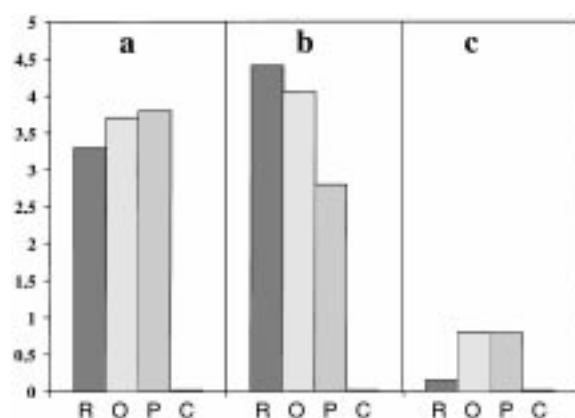


Figure 2. Average disease index of asparagus seedlings, 30 days after inoculation with *F. redolens* (R), *F. oxysporum* (O), or *F. proliferatum* (P). C, noninoculated control. (a) Asparagus cv. Gijnlim grown in tubes with water agar and inoculated with strains CWB1 (*F. redolens*), IPO 97-102 (*F. oxysporum*) or IPO 97-101 (*F. proliferatum*). Root damage was assessed on an ordinal scale from 0 (unaffected) to 5 (completely rotten and hollowed-out; shoots also dead). (b) Asparagus cv. Thielim grown in pots with vermiculite and inoculated with strains IPO 97-13 (*F. redolens*), IPO 97-14 (*F. oxysporum*) or IPO 97-10 (*F. proliferatum*). Root system wounded at inoculation. Shoot damage was assessed on an ordinal scale from 0 (healthy plants) to 5 (dead plants). (c) As in B but root system not wounded at inoculation.

Table 3. Origin of 86 isolates of *F. oxysporum*, *F. redolens* and *F. proliferatum* from asparagus from the Netherlands

	Roots	Spears
<i>F. oxysporum</i>	46	21
<i>F. redolens</i>	3	1
<i>F. proliferatum</i>	11	4

entirely healthy. With a second isolate (IPO 97-13), similar results were obtained in a greenhouse test on cv. Thielim plantlets grown from disinfected seed in vermiculite in 200 ml Fibracan polystyrene pots at 20 °C and inoculated after 20 days with 10 ml conidial suspension ( $10^4$  conidia ml<sup>-1</sup>). After 30 days, shoots had developed disease symptoms with all three *Fusarium* species, provided that the roots had been damaged at inoculation by cutting off part of the roots (Figures 2b and c).

No support was obtained for the hypothesis of Cheah (1986) and Falloon and Tate (1986) that *F. redolens* is specifically associated with spear root, rather than root and crown rot (Table 3). Indeed, Gerlach and Pag (1961) had already reported *F. redolens* from diseased asparagus roots, rhizomes, crowns and stem base lesions. Similarly, UnterEcker (1972) reported *F. redolens* as a root rot pathogen of asparagus. Gordon-Lennox and Gindrat (1987) reported the abundance of *F. redolens* (as *F. oxysporum* var. *redolens*) on necrotic asparagus roots; isolates of *F. redolens* from such roots were all highly pathogenic to asparagus seedlings.

In conclusion, data from our experiments as well as from the literature convincingly show that *F. redolens* is a specialized pathogen of asparagus associated with root rot, crown rot and spear rot. In line with previous descriptions of host-specific forms of *F. redolens*, we propose the following name for the pathogen involved:

#### *Fusarium redolens* Wollenw. f.sp. *asparagi* Baayen

The fungus involved has been reported from Europe (present data), Canada (Gerlach and Pag, 1961), and New Zealand (Cheah, 1986; Falloon and Tate, 1986). Further investigations on strains from asparagus considered to belong to *F. oxysporum* may reveal a widespread occurrence of this fungus throughout asparagus production areas all over the world.

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