

Further identification of races of *Cladosporium fulvum* (*Fulvia fulva*) on tomato originating from the Netherlands, France and Poland

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Accepted 3 February 1989

Abstract

Races of *Cladosporium fulvum*, which can overcome the resistance of the genes Cf₂, Cf₄, Cf₅, Cf₈, Cf₉ and Cf₁₁ have appeared in the Netherlands, France and Poland. Known isolates from the Netherlands and France and three new isolates from Poland have been investigated for the presence of virulence genes using a set of genotypes carrying resistance genes Cf₂ to Cf₁₁. Several Dutch isolates of races, earlier designated as 2.4, 2.4.5 and 2.4.5.9, were found to break down the resistance gene Cf₁₁. These races must therefore be designated as 2.4.11, 2.4.5.11 and 2.4.5.9.11 respectively. In the new Polish isolates virulence genes, overcoming the resistance genes Cf₂, Cf₄, Cf₈, Cf₉ and Cf₁₁ were found. Since all races able to grow on genotypes with Cf₄, could also grow on genotypes carrying Cf₈, it was impossible to discriminate between the genes Cf₄ and Cf₈. These Polish isolates were designated as races 4.11, 2.4.11 and 2.4.9.11. The consequences of the occurrence of these races for tomato breeding are discussed.

Additional keywords: gene-for-gene interaction, leaf mold disease, *Lycopersicon esculentum*, resistance, susceptibility.

Introduction

Cladosporium fulvum (*Fulvia fulva*), the causal agent of leaf mold disease in tomato (*Lycopersicon esculentum* Mill.) has been subject of intensive research for several decades. Many resistance genes have been reported, but new races, overcoming the resistance genes used, have evolved rapidly (Boukema, 1981; Leski, 1977; Laterrot et al., 1985; Laterrot, 1986). These new races are detrimental to the tomato production. Therefore they provoke a continuous need to introduce new resistance genes into commercial cultivars. Still the use of resistant cultivars has reduced the occurrence of *Cladosporium fulvum* in commercial tomato production in the Netherlands to an acceptably low level.

Recently, two new isolates have been obtained from glasshouses for commercial

Table 1. Virulence spectrum of some races of *Cladosporium fulvum* from the Netherlands, France and Poland on tomato genotypes.

Tomato genotypes	Resistance genes	Races of <i>Cladosporium fulvum</i>											
		Netherlands						France					
		2	4	5	2.4	2.4.11	2.4.5	2.4.5.11	2.4.5.9.11	2.5.9	4.11	2.4.11	2.4.9.11
Moneymaker	Cf ₀	S*	S	S	S	S	S	S	S	S	S	S	S
Vetomold	Cf ₂	S	R	R	S	nd	S	nd	S	S	R	S	S
Purdue 135	Cf ₄	R	S	R	S	nd	S	nd	S	R	S	S	S
Vagabond	Cf ₂ , Cf ₄	R	R	R	S	S	S	S	S	R	R	S	S
Ontario 7717	Cf ₅	R	R	S	R	R	S	S	S	S	R	R	R
Ontario 7818	Cf ₆	R	R	R	R	R	R	R	R	R	R	R	R
Ontario 7522	Cf ₈	R	S	R	S	nd	S	nd	S	R	S	S	S
Ontario 7719	Cf ₉	R	R	R	R	R	R	R	S	S	R	R	S
Ontario 7716	Cf ₁₁	R	R	R	R	S	R	S	S	R	S	S	S
Estrella		nd	nd	nd	R	R	S	S	S	nd	nd	R	R
Ostona		nd	nd	nd	R	R	R	R	S	S	nd	R	S
Abunda		nd	nd	nd	R	R	R	R	S	nd	nd	R	?

* S = susceptible, R = resistant, nd = not determined, ? = segregation for susceptible and resistant plants.

tomato production in Poland. A third isolate appeared in a cultivar trial at SHRO, Poland. To identify the virulence genes present in these isolates, they were tested on a set of differentials carrying single resistance genes. For comparison also some races, which are important for tomato breeding in the Netherlands, and 2.5.9, recently isolated in France, were included.

Materials and methods

Tomato genotypes and Cladosporium fulvum isolates. The tomato genotypes were from the IVT tomato collection. The choice of genotypes was based on the presence of one or two known resistance genes for *Cladosporium fulvum* (Kanwar, 1981). The races of *Cladosporium fulvum* 2, 4, 2.4, 2.4.11, 2.4.5, 2.4.5.11, and 2.4.5.9.11 were from IPO. Race 2.5.9 was kindly provided by Dr H. Laterrot, Station d'Amélioration des Plantes Maraîchères, Montfavet, France. The Polish isolates were from SHRO, Poland.

Culture of Cladosporium fulvum. The fungus was grown on potato dextrose agar (PDA) at 22 °C. Cultures were maintained by transferring spores with a droplet of sterile water to fresh PDA. For inoculation large quantities of spores were obtained by growing the fungus in flat bottles of 100 ml with a layer of PDA. These cultures sporulated abundantly in about ten days.

Inoculation and resistance tests. All experiments were carried out at IPO. Seedlings of the tomato genotypes were inoculated by spraying them with a suspension of about 10^6 spores ml^{-1} at the stage of two expanded true leaves. After inoculation, plants were kept at 100% relative humidity under plastic cover for at least two days. They were subsequently incubated in a glasshouse at 20 °C for two more weeks. Artificial light was supplemented and humidity was maintained at >70% by a humidifier. The plants were scored visually in two classes, resistant (R)) and susceptible (S) on the basis of disease development.

Experimental design. In two trials genotypes with the resistance genes Cf₀ through Cf₁₁, except for Cf₁, Cf₃, Cf₇ and Cf₁₀ were screened with single isolates of several races of Dutch, French and Polish origin. In the third trial two, seven and three isolates of races, earlier indicated as 2.4, 2.4.5 and 2.4.5.9 respectively (Boukema, 1981), were compared with some other races on a set of relevant tomato differentials. Also the cultivars Estrella, Ostona and Abunda were included in this trial. All trials were performed in three replicates according to a randomized block design with three plants per plot.

Results

The combined results of the three trials are presented in Table 1. The reaction of the tomato differentials to the races 2, 4, 5 and 2.5.9 were identical with those described earlier (Boukema, 1981; Laterrot, 1986). These interactions once more support the gene-for-gene hypothesis and identify both the resistance genes present in the tomato genotypes and the virulence genes present in the *Cladosporium fulvum* races. In the first two trials, the Dutch isolates, earlier designated as 2.4.5. and 2.4.5.9 (Boukema, 1981), showed odd interactions with 'Ontario 7716', carrying resistance gene Cf₁₁. Bas-

Neth. J. Pl. Path. 95 (1989)

ed on these observations a third trial was set up to screen different isolates of the races, earlier designated as 2.4, 2.4.5 and 2.4.5.9, to determine their interaction with Cf₁₁. The results are included in Table 1: one out of two isolates identified earlier as 2.4, three out of seven isolates known as 2.4.5 and all 2.4.5.9 isolates could also overcome the resistance of gene Cf₁₁.

Two Polish isolates showed a combination of virulence genes, not yet observed before. Based on their differential interaction with tomato genotypes with known Cf genes, the Polish isolates must be designated 4.11, 2.4.11 and 2.4.9.11. The latter had been isolated from cv. Ostona, which was included in the third trial. Based on the differential interactions of 'Ostona' with the races tested, the resistance was found to be governed at least by Cf₉. Cf₆ was the only gene found to confer resistance to all isolates tested. However, sometimes some mycelium was observed, indicating a moderate level of resistance. Responses of Cf₈ and Cf₄ were identical for all isolates tested.

Discussion

The new Polish isolates have the virulence gene 11 in common. One of them was isolated from cv. Ostona, a hybrid cultivar carrying at least Cf₉. In the literature confusion exists about the significance of Cf₁₁. Though Kanwar et al. (1980a) consider it a distinct gene, they hesitate whether it is located on chromosome 9 or 12. Besides, it hardly provided protection against race 12 (= 2.4, Kerr et al., 1980). In the Netherlands attack of 'Ontario 7716' by race 2.4.5 and by 2.4.5.9 but not by race 2 had led to the suggestion that the function of Cf₁₁ might be identical to Cf₄Cf₅ (Boukema, personal communication). The present interactions of several Dutch and Polish strains with Cf₁₁ make clear that Cf₁₁ is not identical to Cf₄Cf₅. So, Cf₁₁ is a unique resistance gene. Consequently, some isolates of races, earlier identified as 2.4 or as 2.4.5 and the 2.4.5.9 isolates, contain the virulence gene 11 and must be designated 2.4.11, 2.4.5.11 and 2.4.5.9.11 respectively.

With respect to Cf₄ and Cf₈ the situation might be comparable. However, based on this and earlier studies there is no reason to distinguish a virulence factor 8. In all cases examined up till now, the reaction of Cf₈ is identical to Cf₄. Yet, linkage studies with morphological genetic markers have positioned both genes on different chromosomes (Kanwar et al., 1980a). However, the conclusive allelic test of Cf₄ with Cf₈ has not been executed. As long as no races are found which discriminate between Cf₄ and Cf₈ the existence of virulence gene 8 is not proven.

It is noteworthy that races show up, carrying a virulence gene of which the corresponding resistance gene in the host population is not present. 'Estrella' was the source of some of the 2.4.5.11 isolates and 'Ostona' of 2.4.9.11, while 2.4.5.9.11 was isolated from a Cf₉ line. The susceptibility of 'Estrella' to 2.4.5 and of 'Ostona' to 2.5.9 indicate that they do not contain Cf₁₁, neither did the Cf₉ line. Concerning 2.4.(11) and 2.4.5.(11), there seems to be an about equal chance that the virulence gene 11 is present or absent. Apparently, races can easily convert into a race carrying virulence gene 11. The opposite, c.q. the reversion into avirulence gene 11, might also occur. Consequently the value of resistance gene Cf₁₁ for practical breeding purposes is very limited.

It is remarkable that new races appear, which combine several existing virulence genes with a new gene. For instance, the new Polish races and the latest appearing Dutch race (2.4.5.9.11) carry a rather complex pattern of virulence genes. As in most cultivars which

are currently grown, only one resistance gene is present, there is no strict need for *Cladosporium fulvum* races to combine several virulence genes. The observation that such complex sets of virulence genes are present in one race, implies that the gain in virulence genes does not cause a decrease of fitness. The consequence for practical tomato breeding is the need to introduce new resistance genes, whenever new races with complex virulence genes appear at a large scale. It has been suggested that the use of tomato cultivars with more than one new resistance gene should slow down the appearance of new races with new virulence genes (Laterrot, 1981). It is essential to state that a combination of two new genes is only effective if both genes are not yet overcome by current isolates.

Though presently the tomato production does not encounter unsurmountable problems caused by *Cladosporium fulvum*, this situation might change. For the time being, Cf₆ still confers resistance to all races available. Other resistance genes are available (Kanwar et al., 1980b; Laterrot, 1981) but the level of resistance, conferred by some of these genes, is rather low.

Samenvatting

Verdere identificatie van fysio's van Cladosporium fulvum (Fulvia fulva) op tomaat afkomstig uit Nederland, Frankrijk en Polen

Fysio's van *Cladosporium fulvum*, die de resistentie-genen Cf₂, Cf₄, Cf₅, Cf₈, Cf₉ en Cf₁₁ kunnen doorbreken, zijn in Nederland, Frankrijk en Polen opgetreden. Met behulp van een groep genotypen, die de resistentie genen Cf₂ tot en met Cf₁₁ dragen, zijn Nederlandse, Franse en enkele nieuwe Poolse isolaten onderzocht op de aanwezigheid van virulentiegenen. Enkele Nederlandse isolaten, eerder aangeduid met 2.4, 2.4.5 en 2.4.5.9, bleken het resistentie-gen Cf₁₁ te kunnen doorbreken. Deze moeten daarom aangeduid worden als respectievelijk 2.4.11, 2.4.5.11 en 2.4.5.9.11. In de nieuwe Poolse isolaten werd virulentie gevonden voor Cf₂, Cf₄, Cf₈, Cf₉ en Cf₁₁. Alle fysio's die op genotypen met Cf₄ konden groeien, groeiden ook op genotypen met Cf₈. Daarom kon geen onderscheid gemaakt worden tussen Cf₄ en Cf₈. De Poolse isolaten behoren tot de fysio's 4.11, 2.4.11 en 2.4.9.11. De gevolgen van het voorkomen van deze fysio's voor de tomateveredeling worden besproken.

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