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Growth and isoenzyme comparison of five isolates of *Venturia inaequalis*

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Observation of the mycelial aspect of five isolates representing the five races of *Venturia inaequalis* indicates that it is possible to differentiate these isolates by their growth pattern and the isolate of race 3 appears to be the most easily distinguishable. Fifteen enzyme systems and non-specific proteins have been analyzed by polyacrylamide and starch gel electrophoresis. Thirteen of the isoenzymes systems and non-specific proteins showed no variation among the five isolates. Esterases isoenzyme patterns allow separation of the isolates since only one isoenzyme is common to all isolates and two others are shared by isolates of race 1 and 5. One carbonic anhydrase isoenzyme was observed to be specific to the isolate of race 2.

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Nous avons observé l'aspect du mycélium de cinq isolats du *Venturia inaequalis*. Des différences ont été constatées entre les différents isolats des races et l'isolat de la race 3 se distingue le plus des autres. Quinze systèmes enzymatiques et les protéines non-spécifiques ont été testés par électrophorèse sur gel de polyacrylamide ou d'amidon. Treize des systèmes enzymatiques ainsi que les protéines non-spécifiques révélés ne montraient pas de variation entre les isolats des races. Le patron isoenzymatique des estérases permet de séparer ces isolats car une seule isoenzyme est commune à tous et deux autres ne se retrouvent que chez les isolats des races 1 et 5. Une isoenzyme de l'anhydrase carbonique est spécifique à l'isolat de la race 2.

Introduction

The apple scab fungus, *Venturia inaequalis* (Cke) Wint., is a major pathogen of apple (*Malus domestica* Borkh.), as it attacks almost all varieties. It infects leaves, flowers and fruits and makes the latter unfit for sale. Palmiter (1934) described *in vitro* variations in morphological and physiological characters of the fungus and Herbst *et al.* (1937) assumed that the morphotypes of this fungus constituted hereditary races. Five races have been identified by their specific virulence on different varieties of apple (Julien and Spangelo 1957; Shay and Williams 1956; Shay *et al.* 1962; Williams and Brown 1968). Race 1 is commonly encountered in Northern America and Europe (Shay and Williams 1956). Race 2 was collected in Southern Dakota and attacks the Dolgo variety, some seedlings of Russian origin (R12740-7A) and the Geneva variety. Race 3 collected in Nova Scotia is virulent against the Geneva variety (Shay and Williams 1956). Race 4 has been isolated from

seedlings R12740-7A that were previously resistant to race 2 (Shay *et al.* 1962). Race 5 appeared simultaneously in France and in England in 1968 and attacks plants possessing the V_m gene for resistance (Williams and Brown 1968).

Polyacrylamide and starch gel electrophoresis have been used widely to classify organisms at various taxonomic levels. Chang *et al.* (1962) applied them to *Neurospora* sp., Clare (1963) to species of *Pythium*, and Durbin (1966) to *Septoria* sp. Differences in the protein bands of two races of *Puccinia graminis* Pers. were found (Macko *et al.* 1967) but Shipton and Fleischmann (1969) were unable to differentiate two races of *Puccinia coronata* Corda on the basis of disc gel electrophoresis of soluble proteins. Howes *et al.* (1982), using two-dimensional isoelectric focusing (IEF) coupled with polyacrylamide gel electrophoresis, found that each of four races of *Puccinia graminis* f. sp. *tritici* Pers. Eriks. & E. Henn. had a unique combination of polypeptides and some of these could be associated with specific fungal genes for virulence. Kim *et al.* (1984) have associated a group of polypeptides with races that are

avirulent or virulent on hosts with specific genes for resistance or susceptibility. Differences in isoenzyme and protein patterns of five isolates of *Cryphonectria cubensis* (Bruner) Hodges were detected by Alfenas *et al.* (1984). Two races of the aggressive strain of *Ophiostoma ulmi* (Buism.) Nannf. were differentiated by Jeng *et al.* (1988) using isoenzyme and protein patterns.

Although electrophoretic techniques have been used extensively in fungal taxonomy, electrophoresis of isoenzymes has not been too much utilized to identify and characterize strains, formae speciales, isolates or races of fungi differing in pathogenicity. A study was initiated to determine the isoenzyme and protein patterns of five isolates representing the five races of *Venturia inaequalis* used in resistance testing and dwarf apple tree selection program by polyacrylamide and starch gel electrophoresis. The purpose of this study was to characterize this material more thoroughly and to have a basis for classifying other isolates.

Materials and methods

Culture procedure. Stock cultures were supplied by Agriculture Canada Research Station, Saint-Jean-sur-Richelieu, Québec. Each race was represented by one isolate. Those isolates were kept under mineral oil in culture tubes maintained at 4°C. They were inoculated on malt extract agar and kept at room temperature (20 ± 2°C). For diametral growth measurements, the inoculum was a standardized disk (5 mm) cut from the perimeter of a 25-day-old colony with a No. 2 cork borer. Two series of four replicates of cultures of each isolate were observed and diametral growth and morphological characters of the cultures were recorded after 23 and 40 days of growth. For dry weight determination, the fungus was grown in liquid medium (10 mL) consisting of malt extract 2%, pH 6.0 (Difco) in square bottles (120 mL) placed on their side. After 20 days of growth at room temperature, the mycelium of 21 bottles was harvested by filtration, rinsed with distilled water, freeze-dried and weighed. A oneway ANOVA analysis was conducted for both types of cultures using the Number

Cruncher Statistical System (NCSS) (Hintze 1986).

Protein extraction. The freeze-dried mycelium was ground in a chilled (4°C) mortar. The extracting solution used for polyacrylamide gel electrophoresis was 0.1 M tris-HCl, pH 6.7 and containing 40% sucrose. For starch gel electrophoresis, the extracting solution was 0.1 M KH₂PO₄ buffer, pH 8.0 and containing 20% sucrose and 20 mg/mL polyvinylpyrrolidone P-40 (Sigma). The slurry was centrifuged at 4°C for 10 min at 600 g and the supernatant was used for electrophoresis.

Electrophoresis. Polyacrylamide gel electrophoresis (PAGE) using the discontinuous buffer system described by Simon (1986) was performed vertically in a simplified apparatus according to Chapel *et al.* (1974) and Raymond (1964). Depending on the enzyme system to be analyzed, two different buffers were used for the starch gel electrophoresis. A quantity of 42.4 g of starch (Electrostarch, Electrostarch Company) were mixed with 400 mL of a 0.18 M tris-EDTA buffer, pH 8.4, diluted five times for the detection of alkaline phosphatase and carbonic anhydrase. For the analysis of the other systems, we used a 0.036 M tris-citrate buffer, pH 7.5, diluted five times. The electrode compartments were filled with the same undiluted buffers. For the analysis of the dehydrogenases, NAD or NADP was added to the gel (35 µmol) before polymerisation and to the electrode compartments (70 µmol). Electrophoresis was carried out horizontally at a constant current of either 35 mA (160 V) for the tris-EDTA buffer and 50 mA (100 V) for the tris-citrate buffer. The electrophoretic separation was achieved within 4 to 6 h at 4°C. The gel slab was covered with an Ice-pak® to prevent warming.

Isoenzymes were visualized by the methods described by Ayala *et al.* (1972), Harris and Hopkinson (1976), Scandalios (1969), and Shaw and Prasad (1970), using acrylamide or starch gel electrophoresis. General proteins were detected following the method of Chrambach *et al.* (1967). Each experiment was replicated at least twice on independent extracts. The relative mobilities (R_f) with reference to the Kohl-

rausch band (acrylamide gels) or the buffer front (starch gels) were recorded.

Results and discussion

Morphological characters. Each of the five isolates presents a peculiar colony according to its margin and aspect of mycelial mat (Table 1) that was maintained over a two-year period of observation. The isolate of race 3 is the most different. It produces woolen-like mycelial threads that radiate and can be noticed as soon as the fungus starts growing. In contrast to isolate of race 1, the isolate of race 2 grows well into the agar and presents a well-defined colony margin and a brown colouration in the surrounding area of the colony. This colouration appears to be the consequence of the production of a melanoprotein-like pigment (Hignett *et al.* 1977). The isolate of race 4 resembles that of race 2. The isolate of race 5 looks like the isolate of race 1 but shows growth rings on the back of the Petri dish. In addition, greyish white sectors replace the usual greyish green colour and almost no conidia are produced. The isolate

of race 5 may have undergone a mutation process since it has been proved by a study on conidial production of the five races that the original stock of race 5 produces conidia (Ross 1974). It is also possible that the race 5 has a particular temperature optimum for sporulation since cultures isolated from different hosts produce spores at higher temperatures while others cannot produce them at temperatures above 20°C (Barakat *et al.* 1982).

Differentiation of each isolate is also possible when grown in liquid media (Table 1). Some isolates produce a mycelial mat on the bottom of the flasks (isolates of races 1, 3 and 4) while others develop small isolated colonies (isolates of races 2 and 5). The mycelium of the isolate of race 3 is not fixed on the bottom of the flask and sets apart in one block. While the colour of the culture medium is pale yellow, it turns dark green when colonized by isolates of races 2 and 3 due to the presence of fungal pigments. After three weeks of growth in the liquid medium, these characteristics fade out and it becomes more difficult to differentiate those isolates.

Table 1. Morphological characters of the five isolates of *Venturia inaequalis* after 23 days of growth on malt agar (a) or 20 days of growth in malt extract broth (b)

Isolate	Culture medium	Morphological characters [§]
1	a	Translucent mycelium growth on perimeter, downy, greyish green.
	b	Coalescent colonies, greyish green, attached to the bottom of the flask.
2	a	Thick mycelium, velvety, greyish dark green.
	b	Isolated colonies, dark green.
3	a	Woolly, dark grey to olive green.
	b	Mycelial mat, green, not attached on the bottom of the flask.
4	a	Thick mycelium, less defined margin than isolate 2. Velvety-downy, brownish green.
	b	Mycelial mat, brownish green.
5	a	Looks like isolate 1, growth rings greyish green and greyish white sectors.
	b	More or less isolated colonies, greyish green.

[§] Adjectives are attributed according to Nobles (1965).

To estimate their growth rate, we measured the diameter of the colonies over a period of 65 days (Fig. 1). The isolate of race 3 showed always the largest diameter and the isolate of race 1 had the smallest one. With one exception, the growth of isolates of races 1 and 3 was always significantly different ($P \leq 0.05$) from all the other isolates using the Fisher and Duncan tests when measured at 15, 25 or 38 days of growth. After 65 days of growth, the mean diametral growth of isolates 1, 3 and 4 were significantly different from all the other isolates ($P \leq 0.05$). The isolate of race 3 produced the highest dry weight while the smallest dry weight was observed for the isolate of race 5 (Fig. 2). Weight differences between all pairs of isolates were significant ($P \leq 0.05$) except between isolates of races 1 and 2. Of all isolates, race 3 was easily distinguishable by the type of mycelial surface as well as by its higher growth rate and dry weight production. However, the other isolates show particular mycelium characteristics that permit them to be readily differentiated after a three-week period on agar.

Electrophoretic analysis. Fifteen enzyme systems were resolved in our study, two of them by PAGE and thirteen in starch gels (Table 2; Fig. 3). In addition, general soluble proteins were also detected by PAGE. Repeated attempts to analyze acid phosphatases (ACP), glutamate-oxalacetate transaminase (GOT) and peroxidases (PX) using both types of gels gave poorly resolved patterns which could not be analyzed with confidence. No variation in the mobility of isozymes among the five *V. inaequalis* isolates were observed for thirteen systems (Fig. 3). Isozyme patterns of glucose-6-phosphate dehydrogenase (G-6-PDH), alcohol dehydrogenase (ADH), phosphoglucose isomerase (PGM), malic enzyme (ME), leucine aminopeptidase (LAP), malate dehydrogenase (MDH), lactic dehydrogenase (LDH), isocitrate dehydrogenase (IDH) and 6-phosphoglucose dehydrogenase (6-PGD) showed one isozyme of similar electrophoretic mobility in the electropherograms of the five *V. inaequalis* isolates.

Invariant patterns were also observed for the two component isozyme patterns of glutamic dehydrogenase (GDH), alkaline phos-

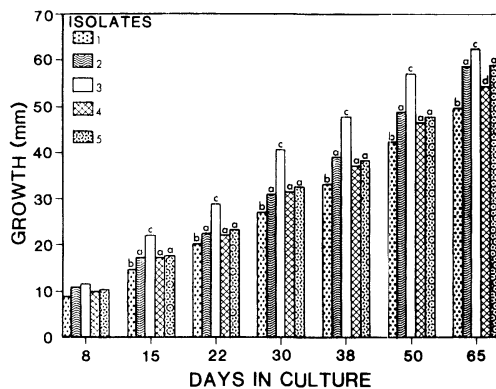


Figure 1. Mean diametral growth of the five isolates of *V. inaequalis* grown on malt extract agar (four replicates). Identical letters indicate means of values not significantly different ($P > 0.05$) according to Duncan's multiple range test.

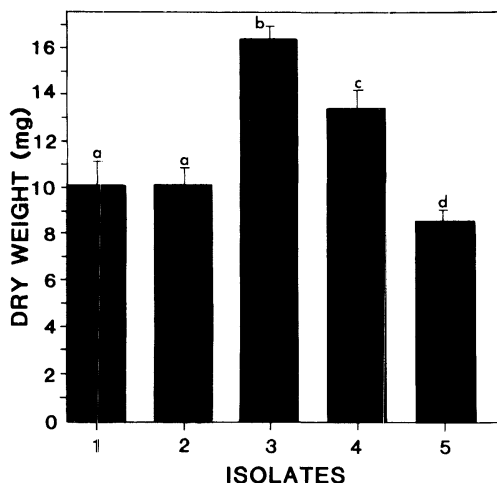


Figure 2. Mean dry weight of the five isolates of *V. inaequalis* grown for 20 days in malt extract broth (21 replicates). Identical letters indicate means of values not significantly different ($P > 0.05$) according to Duncan's multiple range test.

phatase (AKP) and phosphoglucose isomerase (PGI), and for superoxide dismutase (SOD) with a more complex pattern of seven isozymes. General soluble protein (GP) patterns with four well-defined bands were identical for all isolates. Only two enzyme systems, esterase (EST) and carbonic anhydrase (CA) showed variations in the electropherograms that were specific to some of the isolates. The electropherograms of EST analyzed by PAGE revealed two

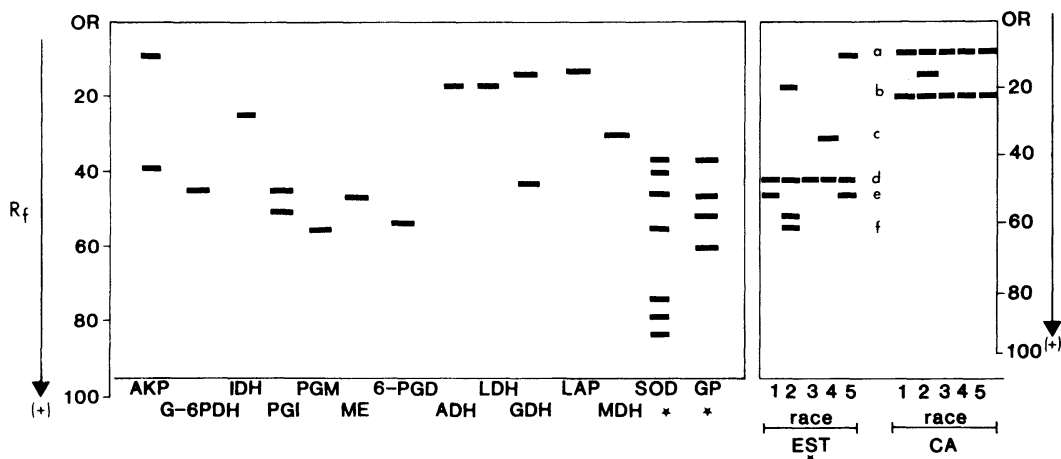
Table 2. Enzyme systems assayed after polyacrylamide (A) and starch (S) gel electrophoresis and staining procedures

Enzyme	Staining procedure	Gel
Esterase (EST)	Scandalios (1969)	A
Superoxyde dismutase (SOD)	Ayala <i>et al.</i> (1972)	A
Acid phosphatase (ACP)	Scandalios (1969)	A
Peroxidase (PX)	Scandalios (1969)	A
Glutamate-oxalacetate transaminase (GOT)	Shaw and Prasad (1970)	A
General proteins (GP)	Chrambach <i>et al.</i> (1967)	A
Carbonic anhydrase (CA)	Harris and Hopkinson (1976)	S
Glucose-6-phosphate dehydrogenase (G-6-PDH)	Harris and Hopkinson (1976)	S
Isocitrate dehydrogenase (IDH)	Harris and Hopkinson (1976)	S
6-phosphogluconate dehydrogenase (6-PGD)	Harris and Hopkinson (1976)	S
Glutamate dehydrogenase (GDH)	Harris and Hopkinson (1976)	S
Alcohol dehydrogenase (ADH)	Harris and Hopkinson (1976)	S
Lactate dehydrogenase (LDH)	Harris and Hopkinson (1976)	S
Malate dehydrogenase (MDH)	Harris and Hopkinson (1976)	S
Malic enzyme (ME)	Harris and Hopkinson (1976)	S
Phosphoglucose isomerase (PGI)	Harris and Hopkinson (1976)	S
Phosphoglucomutase (PGM)	Harris and Hopkinson (1976)	S
Glutamate-oxalacetate transaminase (GOT)	Shaw and Prasad (1970)	S
Leucine aminopeptidase (LAP)	Shaw and Prasad (1970)	S
Alkaline phosphatase (AKP)	Scandalios (1969)	S

(isolate of race 1) to four isozymes (isolate of race 2). Two low R_f isozymes for isolates of race 2 (band b) and 5 (band a) distinguish these isolates from the others (Fig. 3). A well-stained isoenzyme (band e) was only shared by isolates of races 1 and 5 (Fig. 3). Two fast migrating isozymes were present in the isolate of race 2, but their activities were too faint to use them as reliable markers (Fig. 3, band f). Isolates of races 3 and 4 were not easily distinguished although in 60% of the replicated experiments we observed band c in the isolate of race 4 that was never observed in the other four isolates

(Fig. 3). Staining for the enzyme carbonic anhydrase (CA) revealed one major difference between the isolate of race 2 and isolates of the other races. One band was typical of the isolate of race 2.

Burdon *et al.* (1981) also detected differences in the isoenzyme patterns of EST among formae speciales of *P. graminis* pathogenic to different host species. In addition they noticed differences in the isoenzyme patterns of LAP, PGM and dihydrolipoamide reductase. Alfenas *et al.* (1984) working with *C. cubensis*, detected

**Figure 3.** Isoenzyme and general protein banding patterns of *V. inaequalis* as assayed after starch and polyacrylamide* gel electrophoresis. Abbreviations of enzyme systems are given in Table 2.

differences in the isoenzyme patterns of EST, PX, PGM and hexokinase (HK) among five isolates that differ in virulence. Aggressive and non-aggressive isolates of *O. ulmi* were differentiated using isoenzymes of ADH, EST and PX (Bernier *et al.* 1983). Very recently, Jeng *et al.* (1988) reported that isoenzymes of peroxidase, tetrazolium oxidase, and certain protein bands could be used as biochemical markers to distinguish two races of *O. ulmi*. However, formae speciales of *Fusarium oxysporum* Schlecht. could not be distinguished by their esterase and protein patterns (Meyer and Renard 1969).

Also the non specific proteins of all isolates of *C. cubensis* tested by Alfenas *et al.* (1984) displayed nearly the same patterns and neither Shipton and Fleischmann (1969) nor Gill and Powell (1968) could separate races of respectively *P. coronata* f.sp. *avenae* Eriks. and eight races of *Phytophthora fragariae* Hickman on this basis. However, Macko *et al.* (1967) and Jeng and Hubbes (1983) were able to differentiate respectively races of *P. graminis* f. sp. *tritici* and aggressive and non-aggressive isolates of *O. ulmi*. Characterizing races or isolates of fungi by means of protein profiles seems to depend on the organism tested. Our results indicate that only EST isoenzymes could serve as reliable systematic markers for the five isolates of the races of *V. inaequalis* identified by their virulence on differential hosts. It would be desirable to examine the isoenzyme patterns of additional isolates to test this possibility.

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