Phylogenetic aspects of pathogenesis-related (b) proteins

A. CORNU and PATRICIA AHL (Station d'Amélioration des Plantes, Institut National de la Recherche Agronomique, B.V. 1540, 21034 Dijon-Cedex, France)

Biochemical analysis of phylogenetic relationships in plant genera and species has recently been developed as a way for studying evolution in plants. The use of b-proteins for such analyses has several advantages: stable and reproducible protein patterns, a specific variability inherited by simple Mendelian laws, a readily available molecular and genetic approach. Up to now such investigations have only been extensively undertaken in *Nicotiana* species: seven different b-proteins have been recognized in about 15 species belonging to the three main subgenera (1).

Similar genetic analyses have also been made using leaf peroxydase banding patterns (2), and chloroplast-DNA restriction patterns (3). Results from the three methods of analysis show some convergence and broadly confirm the previous classification for species in the genus *Nicotiana* established by Goodspeed (4). The Australian group (subgenus *suaveolentes*) seems to be characterized by the proteins b_1 and b_2 (belonging to the same serological group (5)) and occurrence of b_2 in N. tomentosiformis and N. rustica supports the maternal filiation suggested by the work on chloroplast-DNA patterns. The hypothetical origin of tobacco, (N. tabacum = N. sylvestris \times N. tomentosiformis amphidiploïd) suggested by many workers (cf.1) is also supported by the results of the b-protein analyses; three of the main b-proteins (b_1 , b_2 , and b_3) found in tobacco, probably come from the two hypothetical parents: b_2 from N. tomentosiformis, b_1 and b_3 from the ancestral N. sylvestris.

Although available results and data are still incomplete, it appears from these studies that b-proteins could be usefully employed in phylogenetic investigations.

- (1) Ahl, P., Cornu, A. & Gianinazzi, S., 1982. Soluble proteins as genetic markers in studies of resistance and phylogeny in *Nicotiana*. Phytopathology 72: 80-85.
- (2) Sheen, S.J., 1970. Peroxydases in the genus Nicotiana. Theor. Appl. Gen. 40: 18-25.
- (3) Kung, S.D., Zhu, Y.S. & Shen, G.F., 1982. *Nicotiana* chloroplast genome III. Chloroplast DNA evolution. Theor. Appl. Gen. 61: 73-79.
- (4) Goodspeed, T.H., 1954. The genus Nicotiana. Chronica Botanica Comp., Waltham, Mass., USA, 536 pp.
- (5) Ahl, P., 1983. Aspects génétiques et moléculaires de la résistance (RH) chez les *Nicotiana*. PhD Thesis, University of Geneva, Switzerland.

A new potential for enhancing resistance to tobacco mosaic virus in Nicotiana species

PATRICIA AHL, S. GIANINAZZI and A. CORNU (Station d'Amélioration des Plantes, Institut National de la Recherche Agronomique, B.V. 1540, 21034 Dijon-Cedex, France)

Crosses between *Nicotiana glutinosa* and *N. debneyi* produce hybrids which are highly resistant to tobacco mosaic virus (TMV) and to tobacco necrosis virus (TNV). Healthy plants of these hybrids synthesize a protein (b₁...) which is not present in the healthy parents, but whose appearance can be induced in the parents by infection with