

Limited chloroplast gene transfer via recombination overcomes plastome-genome incompatibility between *Nicotiana tabacum* and *Solanum tuberosum*

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Received 10 August 1988; accepted 12 October 1988

Key words: chloroplast DNA recombination, *Nicotiana tabacum*, plastome-genome interaction, protoplast fusion, restriction enzyme analysis, *Solanum tuberosum*

Abstract

Green cybrids with a new nucleus-chloroplast combination cannot be selected after protoplast fusion in the intersubfamilial *Nicotiana-Solanum* combination. As an approach to overcome the supposed plastome-genome incompatibility, a partial plastome transfer by genetic recombination has been considered. After fusions of protoplasts of a light-sensitive *Nicotiana tabacum* (tobacco) plastome mutant and lethally irradiated protoplasts of wild-type *Solanum tuberosum* (potato), a single green colony was recovered among 2.5×10^4 colonies. The regenerated plants had tobacco-like (although abnormal) morphology, but were normally green, and sensitive to tentoxin, demonstrating chloroplast markers of the potato parent. Restriction enzyme analysis of the chloroplast DNA (cpDNA) revealed recombinant, nonparental patterns. A comparison with physical maps of the parental cpDNA demonstrated the presence of a considerable part of the potato plastome flanked by tobacco-specific regions. This “potacco” plastome proved to be stable in backcross and backfusion experiments, and normally functional in the presence solely of *N. tabacum* nucleus.

Introduction

One of the most stimulating areas of research into cell organization and phylogenesis is the development and mechanism of the delicately concerted interplay between nuclear and organelle genomes [3, 30]. The challenge is to explore its genetic background concealed by a network of pleiotropic nucleus-organelle-environment interactions. One of the approaches that may be adopted is the analysis of plants possessing functional chloroplasts with a plastome derived entirely or partially from a very different species. Both traditional and somatic cell

genetics offer methods for the creation of such cybrids, but there are still major limitations to their application. In genera with biparental organelle inheritance (most thoroughly studied in *Oenothera*) even interspecific crosses may result in various chlorophyll deficiencies, interpreted as different forms of a genome-plastome incompatibility [19, 43]. Protoplast fusion techniques have opened the way to chloroplast transfer between species from distinct genera or tribes [14, 18, 42], but the failure to produce functional *Nicotiana-Solanum* cybrids [42] also suggests a taxonomical barrier to the transfer of entire chloroplast genomes in more remote species

combinations. In the unicellular-uniplastidic green alga *Chlamydomonas*, where sexual mating naturally involves the fusion of the parental chloroplasts, chloroplast DNA recombination has been used extensively for mapping different plastid mutations genetically [33, 12] and physically [21, 22]. In higher plants an obvious barrier to the exchange of plastid genes by sexual crosses is the maternal (cytoplasmic) inheritance of organelles in most species [43, 36, 5]. Furthermore, interorganellar chloroplast DNA recombination appears to be a rare phenomenon, because no spontaneous appearance of genetic recombination of plastids was detectable, even if the plastid markers used allowed visual identification of subgroups of cells with recombinant plastids, either after protoplast fusion [10], or in sexual crosses of species with biparental organelle inheritance [4]. A stringent selection for the recombination of different parental plastid markers in protoplast fusion derived cell cultures did result in the isolation of a single *N. tabacum*-*N. plumbaginifolia* somatic hybrid plant possessing stable recombinant chloroplasts [24].

Biochemical analysis of this plastome demonstrated numerous homologous recombinations all around the genome [24, 9]. This result raised the prospect that disturbances in the cooperation between the nuclear and plastid genomes in remote species combinations might be overcome by a partial plastome transfer via genetic recombination. This approach, whereby a particular mosaic of the parental plastid genes in the recombinant plastome might maintain cooperativity to one parental nucleus, while allowing substantial plastid gene transfer from the other parent, has been attempted between *N. tabacum* and *S. tuberosum*, species from different subfamilies [6] of the family Solanaceae.

Materials and methods

Plant material

Wild-type *Solanum tuberosum* cv. Cardia was maintained as shoot cultures on RM salts [28] with 2% sucrose and 0.7% Bacto-Agar (RM medium) supplemented with choline dichloride (CCC)

(0.3 mg/ml), in light (1500 lx, 16-h day, 26°C). A light-sensitive plastid mutant (LS1) of *Nicotiana tabacum* cv. Turkish Samsun [13] was maintained as shoot cultures on RM medium supplemented with thiamine-HCl (1 mg/ml) and 1-naphthaleneacetic acid (0.1 mg/ml), under low light (400 lx, 16-h day).

Protoplast fusion and cell culture

Protoplasts were isolated from leaves [26], a procedure facilitated in the case of potato by a substantial increase in leaf area of aseptically cultured plants on CCC-containing medium. Those of the appropriate partner were irradiated with a dose of 200 J/kg (0.066 J/kg·s dose rate) of ⁶⁰Co gamma rays [27]. Protoplasts were fused as described previously [26], with the exception of the time of polyethylene glycol treatment (1–2 min). Cultures were grown in liquid medium for five weeks, after which green colonies were isolated in solid medium, and plants were regenerated, as described previously [25, 42].

Chloroplast DNA analysis

Chloroplasts were isolated from leaves of aseptically grown plants in a high ionic strength homogenization buffer as described [2]. CpDNA was purified by phenol-chloroform deproteinization [23], digested with restriction enzymes according to the instructions of the supplier (Boehringer Mannheim GmbH), separated by agarose (0.3% and 1.6%) gel electrophoresis (1.75 V/cm, 18 h), and stained with ethidium bromide.

Results

Production of normal tobacco plants with genetically recombinant chloroplasts

Selection of green colonies following fusions of protoplasts of a light-sensitive *N. tabacum* plastome mutant (LS1) and lethally irradiated protoplasts of wild-type *S. tuberosum* led to the recovery of one cell line among 2.5×10^4 bleached colonies. This green

callus showed apparently no regeneration capability. In a small-scale experiment, therefore, irradiated protoplasts of this callus were again fused with those of the LSI mutant. The single green cell line obtained produced green, tobacco-like shoots, but with an abnormal morphology. Considering that the production of a plant with a completely normal morphology is essential to exclude a possible partial nuclear hybridity, irradiated protoplasts of this fusion derived plant were again backfused to the LSI mutant. Of the 72 green colonies which appeared among 3.0×10^4 bleached ones, several produced shoots identical to those of wild type *N. tabacum*. In the meantime the original green callus, after numerous subcultures on regenerating medium, also produced tobacco-like shoots. Plants from each of the three sequential fusion experiments remained green after transfer to the greenhouse, but the first line had very distorted leaves, the second one had a slightly abnormal morphology, and only the third one chosen for further investigation was completely normal. F₁ progeny of the three lines proved to be sensitive to (bleached by) tentoxin (20 mg/l), as is characteristic of *S. tuberosum*, in contrast to *N. tabacum*, which is naturally resistant to this phytotoxin [8]. This marker of the chloroplasts, together with their normal color, suggested the presence of recombinant chloroplasts in the plants. Backcross ((R × Nt) × Nt) progenies of each of the three regenerates were normal green either on RM medium or in soil (500 seedlings examined in each cross).

Restriction enzyme analysis of the recombinant "potacco" plastome

Restriction enzyme analysis of cpDNA of the plants with each of *Bam* HI, *Pvu* II, *Xho* I (Fig. 1), *Kpn* I, *Pst* I, *Sal* I, *Sma* I (not shown) revealed recombinant, nonparental fragmentation patterns, containing some parent-specific fragments from both tobacco and potato, in addition to nonparental and common fragments. A comparison of the three regenerates and their F₁ progenies (six plants tested) demonstrated identical patterns with *Bam* HI, *Pst* I, *Pvu* II, *Sal* I, *Sma* I, and *Xho* I. No difference was detected between wild-type *N. tabacum*

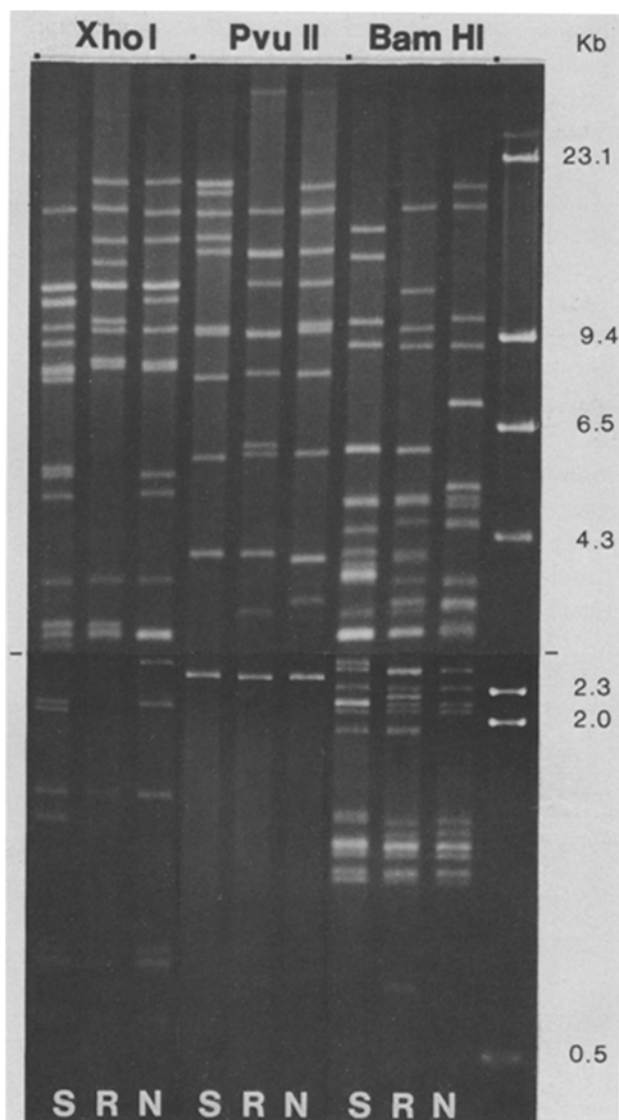


Fig. 1. Restriction endonuclease fragment patterns of chloroplast DNA from *Solanum tuberosum* (S), a tobacco plant with recombinant chloroplasts (R), and *Nicotiana tabacum* (N). On the right a lambda *Hind* III digest is also shown. Fragments larger and smaller than 2.5 kb were separated in 0.3% and 1.6% agarose gels, respectively.

and the LSI mutant with *Bam* HI, *Pvu* II, *Sal* I, and *Xho* I (data not shown). Scoring of individual parent-specific bands in the recombinant cpDNA restriction patterns indicated that only one of the alternative polymorphic parental DNA regions was present, if the region remained parental. A preliminary identification of the position of one or other

of the parental genome parts on the recombinant plastome was feasible due to the availability of physical maps of both *N. tabacum* [37, 11, 34, 16, 41] and *S. tuberosum* [15] chloroplast DNA. The physical map of *Lycopersicon esculentum* cpDNA [31, 32] was also of use because the number and size of tomato cpDNA fragments generated by certain restriction endonucleases are identical with or similar to those of *S. tuberosum* [35]. A comparison of the recombinant cpDNA fragmentation patterns, generated by different enzymes, with the parental patterns and physical maps, demonstrated the presence of numerous potato-specific restriction sites in the inverted repeat region, flanked by tobacco-specific regions (Fig. 2). Further smaller recombination sites in the large single-copy region might also be concluded from a comparison of the proper restriction fragment polymorphisms (Fig. 1).

Discussion

Functional *Nicotiana-Solanum* hybrids with a new nucleus-chloroplast combination could not be selected [42], suggesting an incompatibility of the entire nuclear and plastid genomes in this intersubfamilial combination. *Nicotiana-Solanum* somatic hybrids, however, have a normal green color [38]. During the selection of *N. tabacum* cell lines with recombinant chloroplasts, based on their greening ability, therefore, elimination of the *S. tuberosum* chromosomes was of crucial importance. Ionizing irradiation is effective in the elimination of the irradiated nuclei in protoplast fusion experiments [45, 27], but at least some of the fusion derived plants could be partial hybrids [27]. In the present experiment, regeneration problems and abnormal morphologies in the selected lines, therefore, necessitated repeated backfusions to *N. tabacum* until a completely normal plants was obtained. The lack of appearance of

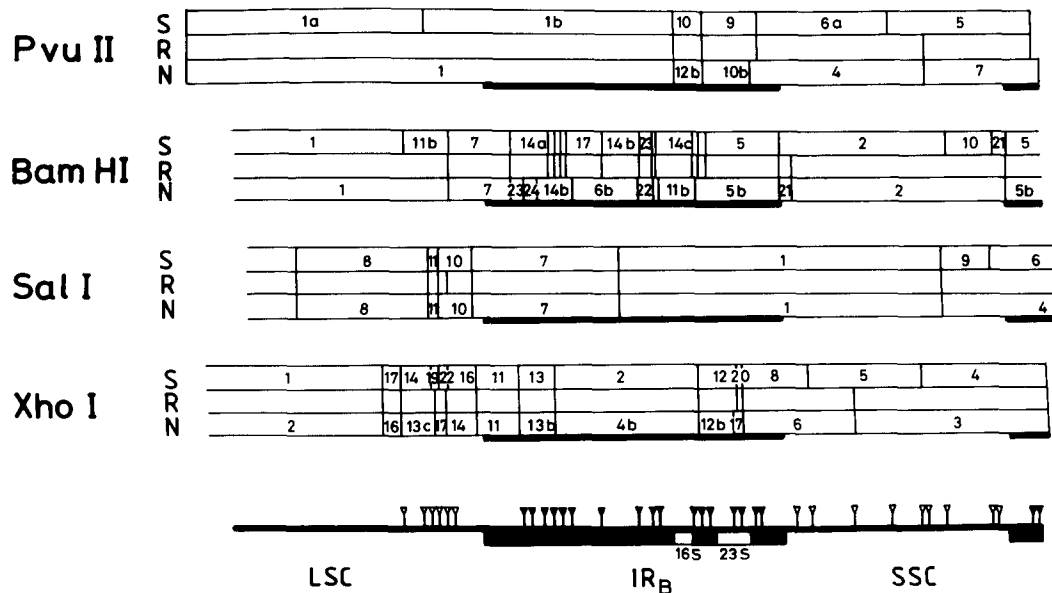


Fig. 2. Comparative restriction endonuclease cleavage site maps of a part of the chloroplast DNA from *Solanum tuberosum* (S), the chloroplast recombinant (R), and *Nicotiana tabacum* (N). The recombinant map was constructed from a comparison of the parental and recombinant restriction fragment patterns with physical maps of tobacco [37, 16, 41], potato [15], and tomato [31, 32, 35] cpDNA. Numbering of the fragments follows the original publications. The inverted repeat regions are marked by thick lines. A summation of the parent-specific restriction sites from tobacco (open triangles) and potato (solid triangles), present in the recombinant plastome, is marked on a simplified map, showing (from left to right) a part of the large single-copy region (LSC), the left inverted repeat (IR_B) with the 16S and 23S rRNA genes, and the full small single-copy region (SSC) [40, 41]. The distribution of the parent-specific restriction sites was also supported by the use of *Kpn* I, *Pst* I, and *Sma* I fragmentation patterns (data not shown).

any kind of chlorophyll deficiencies in regenerates derived from the backfusions, and progenies obtained from the backcrosses, strongly indicated that the abnormality of the first and second regenerates was due to their polyploid-aneuploid chromosome numbers, induced by the cell culture itself [1, 20]. Using the amphidiploid, and therefore relatively unstable *N. tabacum* [39], similar problems have also been reported in other protoplast fusion experiments [42, 7]. In the LS1 plastid mutant no revertants have been found in our and other laboratories [13]. Green color and tentoxin sensitivity of the normal tobacco regenerate was, therefore, a strong indication of the recombinant nature of its chloroplasts. In contrast to the case of the intrageneric *Nicotiana* chloroplast recombinant [24], each of the enzymes used revealed a restriction fragmentation pattern markedly different from those of the parents, which was to be anticipated considering the much wider taxonomical distance [44]. The absence of any variation in the patterns of the vegetative, somatic, or sexual derivatives of the originally selected cell line demonstrates the stability of the recombinant plastome, and the termination of the intraorganellar DNA and intracellular organelle segregation, at the end of the selection period. A preliminary view of the recombinant DNA suggests a homologous recombination between the parental DNAs. It seems that the major step in the formation of the recombinant plastome was an exchange of (at least most of) the inverted repeat region of tobacco with that of potato, which is not surprising considering its high intramolecular recombination activity [29, 17].

To clarify the mechanism of the recombination, to identify all of the recombination sites, and to explore the plastome for possible smaller deletions, insertions, or inversions, fine mapping and sequencing of numerous cloned fragments will be necessary. With respect to the main approach in this work, however, the motivation for a detailed biochemical analysis is frustrated by the great number of genes present from both species, which prevents the localization of the plastid markers (chlorophyll deficiency and tentoxin resistance) used in the selection, and the identification of the genes essential in the cooperation with the nucleus. In this "potacco" plastome it is likely that a particular combination of tobacco genes, crucial

in the interaction with the nucleus, and potato genes, neutral from this viewpoint, but complementing the plastome deficiency, ensure functionality of the chloroplast genome in the presence of a tobacco nucleus. Results presented here demonstrate that chloroplast DNA recombination, brought to light by the power of cell culture selection, may be considered as an efficient tool to achieve chloroplast gene transfer between somatically/sexually incompatible species.

Acknowledgements

The authors thank Irute Meshkiene, Éva Horváth, and Mihály Kis for valuable technical suggestions, Erika Veres for her excellent technical assistance, Philip Dix for helpful discussion of the manuscript, and Béla Dusha for photography. The work was supported by an OKKFT/Tt grant No. 4413.

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