

Agrobacterium-mediated transformation of thin cell layer explants from *Brassica napus* L.*

P. J. Charest¹, L. A. Holbrook²**, J. Gabard², V. N. Iyer¹ and B. L. Miki²***

¹ Department of Biology, Carleton University, Ottawa, Ontario K1S 5B6, Canada

² Plant Research Centre, Agriculture Canada, Ottawa, Ontario K1A 0C6, Canada

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Summary. Agrobacterium-mediated transformation of thin cell layer explants (Klimaszewska and Keller 1985) yielded large numbers of transgenic plants of a major Canadian rapeseed cultivar Brassica napus ssp. oleifera cv Westar. The morphology and fertility of these plants were indistinguishable from controls. The Ti plasmid vector, pGV3850 (Zambryski et al. 1983) was used as a cis vector and as a helper plasmid for the binary vector pBin19 (Bevan 1984). Selectable marker genes that conferred resistance to high levels of kanamycin (Km) on Nicotiana tabacum were less efficient in the selection of transgenic B. napus. At low levels of Km (15 µg/ml) large numbers of transgenic plants (50%) were identified among the regenerants by nopaline synthase activity and several of these were confirmed by Southern blot analyses. Only a small number were resistant to higher levels of Km (80 µg/ml). Preliminary analyses indicated that resistance to Km was transmitted to the selfed progeny. Chimeric chloramphenicol acetyl transferase genes were ineffective biochemical markers in transgenic B. napus.

Key words: Brassica napus – Rapeseed – Transformation – Agrobacterium tumefaciens – Ti plasmid

Introduction

The genus *Brassica* encompasses a wide range of economically-important crop species including rapeseed.

Recent advances in the cell and tissue culture technologies for rapeseed have enabled the genetic manipulation of the cytoplasmic genomes (Pelletier et al. 1983; Yarrow et al. 1986). Continued advances using genetically-engineered nuclear genes depends on the development of methods for producing transgenic plants. *Brassica* species in general are excellent hosts for transformation by Ti and Ri plasmids (Holbrook and Miki 1985; Mathews et al. 1985, 1986; Tanaka et al. 1985; Ooms et al. 1985). Recently, fertile, transgenic *B. napus* have been generated with the Ri plasmid of *A. rhizogenes;* however, these plants expressed the morphological abnormalities and reduced fertility imposed by the Ri T DNA (Guerche et al. 1987).

In this report, the transformation of a major Canadian rapeseed cultivar, *B. napus* ssp. *oleifera* cv. Westar with *A. tumefaciens* strains harboring disarmed Ti plasmid vectors is demonstrated. Thin cell layer explants which yielded consistently high frequencies of plant regeneration with cultivar Westar (Klimaszewska and Keller 1985), were adopted as hosts for transformation. The major obstacle in the regeneration of transgenic plants was the hypersensitive response of the explants to the *A. tumefaciens* strains. Once conditions were determined that overcame this problem, transgenic plants that were indistinguishable from controls were recovered.

Materials and methods

Plants, cultures and strains

Seeds of *B. napus* ssp. *oleifera* cv Westar were supplied by R. K. Downey, Agriculture Canada (Saskatoon, Canada). *Brassica napus*, Jet Neuf, cell suspension cultures were obtained from W. Keller, Agriculture Canada (Ottawa, Canada).

^{*} Contribution No. 1092 Plant Research Centre, Ontario, Canada

^{**} Present address: Biotechnica International of Canada Inc., 170-6815 8 Street NE, Calgary, Alberta T2E 7H7, Canada *** To whom correspondence should be addressed

The A. tumefaciens strains used in this study are described in Table 1. The chimeric genes nos-nptII-ocs (5'region-coding sequence-3' eno), nos-CAT-nos, 35S-CAT-nos (Hain et al. 1985; DeBlock et al. 1984; M. Bevan, unpublished) were evaluated as selectable markers. The nos-CAT-nos and nosnptII-ocs genes were cointegrated with the T DNA of pGV3850 using the intermediate vectors pNCAT4 (DeBlock et al. 1984) and pLGV1103neo (Hain et al. 1985) to form the two cis vectors pGV3850: :pNCAT4 and pGV3850: : pLGV1103 neo. The 35S-CAT-nos gene was inserted into the binary vector pBin19 (Bevan 1984) by M. Bevan to form pBCAT1 (unpublished) and this vector was maintained in A. tumefaciens with pGV3850 acting as the helper plasmid. Vector pBCAT1 also carried a chimeric nos-nptII-nos gene.

Transformation of thin cell layers

Thin cell layer explants used in cocultivation studies with A. tumefaciens were prepared according to the specific conditions determined by Klimaszewska and Keller (1985). Modifications to the tissue culture procedures were made to compensate for the use of the explants in cocultivation experiments. Prior to inoculation with bacteria, B. napus explants were preconditioned by culture over a feeder layer of B. napus, Jet Neuf, cells from suspension cultures using the double filter paper technique (Horsch and Jones 1980). After one to four days, explants were dipped in an overnight bacterial culture grown in LB medium and diluted with distilled sterile H₂O or media. The explants were blotted on filter paper and cocultivated for one to three days. To inhibit bacterial growth, the explants were transferred to media solidified in 0.7% agarose (Seakem ME, FMC) to which 0.5 mg/ml carbenicillin (Ayerst Labs, Montreal, Canada), 0.5 mg/ml Clavulin 250 (Beecham Labs Inc., Montreal, Canada), 0.5 mg/ml cefotaxime (Terochem Lab. Ltd., Mississauga, Canada) or 0.5 mg/ml Claforan (Roussel Canada, Montreal, Canada) was added.

Selection conditions

At various times, the explants were transferred to media that also contained 15 μ g/ml kanamycin (Km) or 7 μ g/ml chloramphenicol (Cm) and subcultured approximately every two weeks. Well-developed shoots were transferred to B₅ media lacking growth regulators to induce roots in the presence of Km or Cm and plantlets were transferred to peat pellets prior to soil (Klimaszewska and Keller 1985). The effect of Km and Cm on growth and differentiation was examined. Explants were transformed with an *A. tumefaciens* strain carrying only pGV3850 (GV3101, Table 1) and cultured on media at six different antibiotic concentrations. A minimum of 12 explants were cultured at each concentration. The fresh weights measured after five weeks and the average fresh weight was expressed as a percentage of the average fresh weight of tissues growing in the absence of Km or Cm.

Opine synthase analyses

Nopaline synthase and octopine synthase activities were analyzed by precursor labelling of nopaline or octopine with [¹⁴C]arginine, followed by paper electrophoresis and fluorography (Holbrook et al. 1986). For each plant examined, approximately 10 randomly selected tissue samples were analyzed.

Southern blot analyses

Plant DNA was isolated from leaves by methods described by Lichtenstein and Draper (1985). Southern blot analyses were performed according to Bernatsky and Tanksley (1986).

Inheritance of antibiotic resistance

Regenerated plants were self-pollinated. The seeds were sterilized with 0.2% (w/v) mercuric chloride, 20–25 min and 6% (w/v) sodium hypochlorite 20–30 min. The seeds were rinsed in sterile distilled water and placed on MS medium with B_5 vitamins, 1.0% sucrose, 0.6% agarose. Resistance or sensitivity to 20 µg/ml Km or 40 µg/ml Cm was scored after five to six weeks.

Results

Transformation of thin cell layer explants

Brassica napus thin cell layer explants (Klimaszewska and Keller 1985) were evaluated as hosts for Agrobacterium-mediated transformation. The uninoculated explants were highly regenerable in culture producing on average, five shoots per explant within three to four weeks. Exposure to A. tumefaciens strains following the

Table 1. Agrobacterium tumefaciens strains and plasmids used for the transformation of Brassica napus

Strain	Chromosomal background	Ti plasmid ª	Intermediate vector ^b	Binary vector ^c	Marker genes
1. C58	C58	pTiC58	_	_	nos (aut) ^d
2. GV3101	C58	pGV3850	-	_	nos
3.		1	pNCAT4	_	nos, nos-CAT-nos
4.			pLGV1103neo	_	nos, nos-nptII-ocs
5.				pAL1050	$ocs, (aut)^{\frac{1}{d}}, (nos)^{e}$
6.			-	pBCAT1	nos-nptII-nos, 35S-CAT-nos, (nos) ^e
7. B6S3	B6S3	pTiB6S3	-	-	ocs, (aut) ^d

^a pTiC58 (Holsters et al. 1980); pGV3850 (Zambryski et al. 1983); pTiB6S3 (Petit and Tempe 1978)

^b pNCAT4 (DeBlock et al. 1984); pLGV1103neo (Hain et al. 1985)

^c pAL1050 (Hoekema et al. 1983); pBCAT1 (M. Bevan, unpublished)

^d (aut). In culture, growth in the absence of growth regulators was used as a selectable marker for T-DNA of natural Ti plasmids ^e (nos). In strains harbouring binary vectors, nopaline synthase activity was used as a marker for transformation by T-DNA of pGV3850

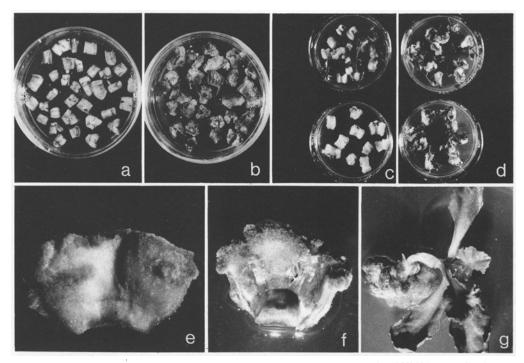


Fig. 1a-g. Thin cell layer explants cocultivated with *A. tumefaciens* strain GV3101. a, e Explants prepared according to Klimaszewska and Keller (1985) and cocultivated using the leaf disc transformation procedure (Horsch et al. 1985) were chlorotic and brown after three weeks. b, f Viable explants with green meristematic regions after three weeks were recovered using the modified conditions for inoculation and culture. d, g Shoot differentiation was achieved by six weeks. c The frequency of shoot differentiation was reduced on media solidified in 0.8% agar

Table 2. Transformation rates of thin cell layer explants of B. napus^a

Strain	Plasmids	Precond	lition-cocultivation ^b	
		0-2	4-2	4-1
C58	pTiC58	50	100	16
B6S3	pTiB6S3	0	48	0
GV3101	pGV3850, pAL1050	N.T.	79	N.T.

^a Transformation rates were measured as the percentage of infected explants that demonstrated nopaline synthase (pTiC58) or octopine synthase (pTiB6S3, pAL1050) activities. Measurements were based on the analyses of at least 15 explants

^b The explants were cocultivated with the \hat{A} . tumefaciens strains for two days without preconditioning (0-2) of after preconditioning for four days (4-2). Explants were also cocultivated for 1 day after preconditioning for four days (4-1) N.T. = not tested

procedure for leaf disc transformation (Horsch et al. 1985) induced a severe hypersensitive response that completely prohibited the recovery of viable tissues (Fig. 1 a, e). The procedures for inoculation and culture of explants were therefore modified to minimize tissue necrosis (Fig. 1 b, d, f, g) and to enhance explant transformation rates (Table 2).

The inoculation of the explants with the bacterial strains was a critical step in the recovery of healthy tissues. With suspensions of 10^9 bacteria per ml, inoculations could not exceed 30s because rapid bacterial growth during cocultivation could not be inhibited by carbenicillin, cefotaxime, Claforan or Clavulin 250 at 500 µg/ml. Shorter inoculations permitted cocultivation periods up to two days and subsequent inhibition of bacterial growth by carbenicillin or Clavulin 250. Cefotaxime and Claforan were toxic to the explants under these conditions of culture. In the recovered tissues necrosis was still evident and the frequency of shoot differentiation was suppressed to about 0.6 shoots per explant (Fig. 1c).

A dramatic reversal of tissue necrosis was achieved by replacing agar in the media with 0.7% agarose (Fig. 1 b, d, f, g). Shoot differentiation was restored to levels of three to five shoots per explant (Fig. 1 d, g); however, this was delayed by one to two weeks relative to uninoculated controls. Preconditioning of explants prior to inoculation with feeder layers of *B. napus* cell suspension cultures did not alter these results significantly but appeared to be associated with greater consistency among experiments in the recovery of viable tissues. Under these improved culture conditions, any of the antibiotics carbenicillin, cefotaxime, Claforan or Clavulin 250 were effective in the control of bacterial growth without detriment to plant tissue growth and differentiation.

Explant transformation rates were assessed by analyses of opine synthase activities in tissues transformed with pTiC58 or pTiB6S3 and selected on media lacking growth regulators. With both strains, a four-day preconditioning period and a two-day cocultivation were effective (Table 2). Longer durations of preconditioning had not been examined. Longer cocultivation resulted in uninhibited bacterial growth. Transformation rates for pTiC58 were consistently high and for pTiB6S3 were always much lower (Table 2). The same conditions applied to a strain harboring the disarmed Ti plasmid vector pGV3850 and the binary T-DNA vector pAL1050 yielded high transformation rates for pAL1050 (Table 2). A large proportion (68%) of these were also transformed with pGV3850 revealed by the presence of both octopine synthase and nopaline synthase activities.

Selection and regeneration of transformants

The effect of the antibiotics Km and Cm on the growth and differentiation of thin cell layer explants is illustrated in Fig. 2. These profiles were determined with explants after cocultivation with a strain carrying only pGV3850 (GV3101, Table 1). At 35 µg/ml Km, shoot differentiation was completely abolished and tissue growth was significantly retarded (Fig. 2). Following cocultivation with strains carrying pGV3850:: pLGV1103 neo or pGV3850 and pBCATl, shoots did not differentiate in the presence of 35 µg/ml Km despite repeated attempts. The same strains were used to recover tobacco shoots resistant to 100 µg/ml Km (data not shown). At 15 µg/ml Km many shoots differentiated but several were gradually lost at later stages of growth and differentiation (Fig. 3a, b). Typically, these shoots became chlorotic and would not form roots before dying (Fig. 3b). Many that survived and formed roots in the presence of 15 µg/ml Km after one to two months (Fig. 2b) were successfully transferred to soil after two to three months. Approximately one-half of the plants displayed nopaline synthase activity following transformation with pGV3850:: pLGV1103 neo. Most of them displayed activity after transformation with pGV3850 and pBCATI; however, nopaline synthase activity in these plants reflected transformation by the T-DNA of pGV3850 whereas the nos-nptII-nos and 35S-CAT-nos genes are carried by pBCATl. A summary of these experiments is presented in Table 3.

Chloramphenicol at levels above 20 μ g/ml inhibited explant growth and differentiation; however, resistant shoots could not be recovered at these levels after

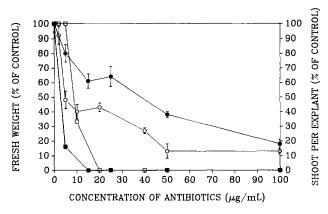


Fig. 2. Influence of Km and Cm on the growth and differentiation of thin cell layer explants. At least 12 explants (fresh weight 10 ± 2 mg) cocultivated with strain GV3101 were incubated on media with antibiotics at each of the concentrations indicated (Km $\bullet - \bullet$, $\blacksquare - \blacksquare$; Cm $\circ - \circ$, $\Box - \Box$). The average fresh weight after five weeks ($\bullet - \bullet$, $\circ - \circ$) were expressed as a percentage of the average fresh weight of explants incubated without antibiotics (601 ± 32 mg). Bars indicate standard errors. The frequency of shoot differentiation ($\blacksquare - \blacksquare$, $\Box - \Box$) was expressed as a percentage of the frequency (shoots per explant) measured for explants incubated without antibiotics

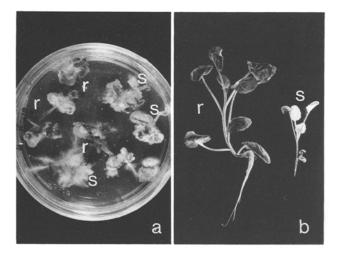


Fig. 3a, b. Selection of transformed shoots and plantlets on Km. Thin cell layer explants were cocultivated with A. tumefaciens harbouring pGV3850::pLGV1103neo and cultured on 15 μ g/ml Km. a Shoot differentiation occurred in the presence of Km; however, many shoots became chlorotic (s) and died. b The survivors (r) were excised and roots were induced in the presence of Km. Many of them did not form roots. They became chlorotic and eventually died (s). Others formed roots in the presence of Km (r), and were transferred to soil

transformation with pGV3850: :pNCAT4 or pBCAT1. Reduction of Cm to 7 μ g/ml yielded many shoots but tissues transformed with pGV3850 alone also yielded shoots at this level despite a modest suppression of tissue growth (Fig. 2). About one half of the plants recovered following tansformation by pGV3850: :

Plasmids Selec-Ex-Resis-Plant-Nopation^a plants tant lets line inocushoots^b in soil synlated thase^d pGV3850:: 40/80 Km 600 26 12 pLGV1103neo pGV3850::pNCAT4 Cm 600 30/70 19 10 pGV3850, pBCAT1° Km, 200 10/406 5 Cm

Table 3. Recovery of transformed B. napus plants

^a Antibiotic selections were performed at 15 μg/mL Km or 7 μg/mL Cm

^b The resistant shoots are indicated as a proportion of the total shoots that differentiate in the presence of low levels of antibiotics

^c Explants transformed with pBCAT1 were subjected to selection on Km or Cm. Each of the selections yielded three plants. These were analyzed for nopaline synthase activity which was encoded by pGV3850

^d Nopaline synthase activity was determined for plants established in soil. At least 10 samples were taken from each plant for analyses because of variability that existed within inidividual plants

Table 4. Inheritance of Km^r in self-pollinated transgenic *B. napus* plants

Plasmids	Plant	Km ^r : Km ^{s, a}
pGV3850::pLGV1103neo	NEO4A	9:6
1 1	NEO12A	8:5
	NEO8B	7:4
	NEO12B	12:4
	Control	0:25
pGV3850, pBCAT1	BCAT1B	7:8
	Control	0:25

* Seedlings were germinated in media with $20 \,\mu\text{g/ml}$ Km and resistance was scored after five to six weeks

pNCAT4 were associated with nopaline synthase activity (Table 3).

Several of the plants potentially transformed with pBCAT1 displayed nopaline synthase activity encoded by pGV3850 (Table 3). CAT activity as a direct biochemical marker for transformation was impossible to assess because all of the tissues were associated with very high levels of endogenous CAT activity (data not shown).

Analyses of transformants

All of the plants that displayed nopaline synthase activity flowered and set seed within five to six months. In morphology and fertility, these plants were indistinguishable from control plants derived from seeds and grown under identical conditions (Fig. 4).

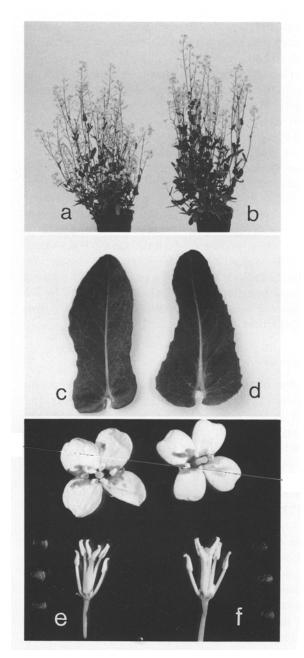


Fig. 4. Comparison of regenerated transgenic plants with normal *B. napus* cv. Westar plants. a Transgenic plant NEO8B transformed with pGV3850::pLGV1103neo. c, e leaf; flower, floral parts, seeds derived from NEO8B. b Normal *B. napus* cv. Westar plants grown from seed. d, f Leaf; flower, floral parts, seeds derived from normal plants

Southern blot analyses confirmed the integration of vector DNA at the right T-DNA border in six out of seven plants tested. With a fragment of the *nos* gene as the hybridization probe, two separate integration events were revealed for a representative plant (NEO12B) transformed with pGV3850: :pLGV1103 neo and another representative plant (NCAT7B) transformed with

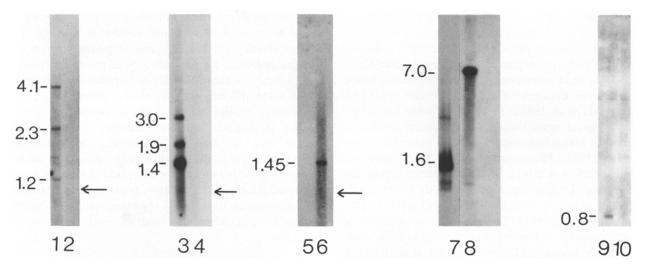


Fig. 5. Southern blot analyses of transgenic plants. The arrows indicate the minimum fragment size that would encompass the T DNA-plant DNA junction at the right border of pGV3850 following digestion with BamH1 and HindIII and hybridization with the BamH1-SstII fragment from the HindIII 23 fragment of pTiC58 as the probe. DNA from transgenic plant NCAT7B transformed with pGV3850::pNCAT4 was digested with BamH1 and HindIII (*lane 1*). The 1.2 kb BamH1-HindIII fragment of pNCAT4 is indicated. Two T DNA-plant junction fragments of 4.1 kb and 2.3 kb were revealed. DNA from transgenic plant NEO12B transformed with pGV3850::pLGV1103neo was digested with PstI, BamH1 and HindIII (*lane 3*). Two T DNA-plant DNA junction fragments of 3.0 kb and 1.9 kb were revealed in addition to the 1.4 kb PstI fragment of pLGV1103neo. DNA from transgenic plant BCAT1B transformed with pGV3850, pBCAT1 was digested with BamH1 and HindIII (*lane 6*). A single T DNA-plant DNA junction fragment of 1.45 kb was revealed. Comparable bands were not found in DNA from untransformed control plants (*lanes 2, 4, 5*). The 1.6 kb PstI fragment of pLGV1103neo was used as the probe to reveal the same fragment which contains the coding region of *npt1I* was also demonstrated in a 7.0 kb fragment from DNA of transgenic plant BCAT1B digested with PstI (*lane 8*). The 0.8 kb BamH1 fragment of pNCAT4 was used as the probe to reveal the same fragment of pNCAT4 was used as the probe to reveal the same fragment of pNCAT4 was used as the probe to reveal the same fragment of pNCAT4 was used as the probe to reveal the same fragment of pNCAT4 was used as the probe to reveal the same fragment of pNCAT4 was used as the probe to reveal the same fragment of pNCAT4 was used as the probe to reveal the same fragment of pNCAT4 was used as the probe to reveal the same fragment of pNCAT4 was used as the probe to reveal the same fragment of pNCAT4 was used as the probe to reveal the same fragment of pNCAT4 was used as the probe to reveal the same fragment of pNCAT4 was used a

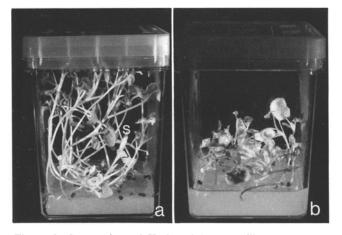


Fig. 6a, b. Segregation of Km^s and Km^T seedlings. a Seeds from a self-pollinated transgenic plant NEO12B and b control *B. napus* plants were germinated in the presence of $20 \,\mu\text{g/}$ ml Km. Sensitive seedlings (s) were chlorotic and growth was inhibited. Resistant seedlings were green and displayed normal growth characteristics

pGV3850: :pNCAT4 (Fig. 5). A single integration event is shown for another plant (BCAT1B) transformed with a strain carrying pBCAT1 and pGV3850 (Fig. 5). The presence of the coding region of the *npt11* gene was confirmed in both NEO12B and BCAT1B and the coding region of the CAT gene was confirmed in NCAT7B (Fig. 5).

To assess the level of antibiotics that could be tolerated, shoots were cloned from the axillary buds of plants selected on 15 µg/ml Km that displayed nopaline synthase activity encoded by pGV3850: : pLGV1103 neo. Shoots from each were subcultured on media with 20, 40, or 80 µg/ml Km. In one transformation experiment, 13 plants were sensitive to 20 µg/ml Km. One plant was tolerant to 40 µg/ml Km but not 80 µg/ml. Another plant was resistant to all of the levels tested. Similar experiments, with eight plants transformed with pGV3850: :pNCAT4 were subjected to various levels of Cm; however, the results were inconclusive due to the inherent and variable tolerance of differentiated *B. napus* shoots to this antibiotic.

Preliminary genetic analyses were performed on a limited number of plants. Seedlings from control plants that germinated in the presence of 20 μ g/ml Km were uniformly chlorotic and severely inhibited in growth (Fig. 6b). Among the progeny of self-pollinated plants that were transformed with pGV3850: :pLGV1103 neo or pBCAT1 resistance to 20 μ g/ml was evident (Fig. 6a).

Discussion

Transgenic B. napus plants were efficiently regenerated from thin cell layer explants (Klimaszewska and Keller 1985) that were transformed by A. tumefaciens strains carrying the disarmed Ti plasmid vector, pGV3850 (Zambryski et al. 1983). These plants were free of the morphological abnormalities and reduced fertility of transgenic plants transformed with Ri plasmids (Guerche et al. 1987). The vector pGV3850, which is derived from pTiC58, was chosen because an earlier survey had shown that B. napus was most efficiently infected with nopaline plasmids such as pTiC58 and pTiT37 (Holbrook and Miki 1985). The virulence of pGV3850 in association with B. napus was confirmed with the binary T DNA vector, pAL1050(Hoekema et al. 1983). It was employed effectively both as a cis vector and as a helper plasmid for the binary vector pBin19 (Bevan 1984).

A survey of methods described for the regeneration of *Brassica* plants from tissue explants (Kartha et al. 1974; Margara and Leydecker 1978; Stringam 1979; Dunwell 1981; Lazzeri and Dunwell 1984a, b; Klimaszewska and Keller 1985) revealed that thin cell layer explants (Klimaszewska and Keller 1985) were particularly regenerable and well suited for the experimental manipulations involved in transformation by *A. tumefaciens*. A major obstacle was the severe hypersensitive response of the tissues to the *A. tumefaciens* strains and the associated loss of morphogenesis. This problem was overcome by determining the optimal conditions for inoculating the explants and by modifying the culture conditions to enhance tissue survival.

Large numbers of transgenic plants have been produced with the methods described in this study. Nopaline synthase activity (Holbrook et al. 1986) was a reliable biochemical marker for identifying them. Chloramphenicol acetyl transferase activity could not be used because of the high levels of endogenous activity found in all of the tissues examined. A similar finding has recently been reported (Balazs and Bonneville 1987). Furthermore, chloramphenicol was a poor selective agent for differentiated shoots and seedlings because of the inherent and variable tolerance of these tissues to this antibiotic. Despite this, a large number of plants transformed with chimeric CAT genes have been recovered and verified by Southern blot analyses. As selectable marker genes nos-nptII-nos (Bevan 1984) and nos-nptII-ocs (Hain et al. 1985) did not provide selection of transformed B. napus shoots that could differentiate fully in the presence of high levels of Km; however, gradual selection at low levels of Km (15 µg/ml) yielded a high proportion of transformants among the regenerated plants (50%). Only a small number of these were also resistant to high levels of Km (80 μ g/ml). Attempts to select for these directly at high levels of Km were not successful. Clearly, the transformation of thin cell layer explants was an efficient approach for the recovery of transgenic plants but further research is needed to improve the efficiency of selection and the selectable marker genes.

Westar is the major *B. napus* rapeseed cultivar grown in Canada. For the introduction of isolated genes into this crop, the procedures described are effective. Subsequent breeding with transgenic plants should not be obstructed by decreased fertility or gross morphological abnormalities; however, the stability and segregation of the marker genes must be evaluated through detailed genetic analysis which is now in progress.

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