An Improved Technique for High Resolution Mitotic Chromosome Studies in Solanum

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Abstract. An improved method is described for the isolation of potato metaphase chromosomes for karyotypic and cytogenetic studies. Root tips from diploid Mexican species, Solanum pinnatisectum (2n = 2x = 24) and tetraploid cultivated S. tuberosum (2n = 4x = 48) were given four different pretreatments. The synthetic pyrethroid, Ambush, was the most stable and effective pretreatment reagent, providing the highest percentage of mitotic chromosomes at metaphase and the best spread of countable chromosomes for cytogenetic studies. Compared with an Ambush pretreatment at concentrations of 100–400 ppm, 1 to 10 ppm Ambush produced more easily distinguished chromosomes, which can be useful for comprehensive observation and karyotype analysis in both 2x and 4x potato species. This improved technique for examining mitotic chromosomes will be helpful in describing karyotypes, characterization of new hybrids, and identifying chromosome structural changes that are important in breeding schemes.

In plants, cytogenetic analysis has been widely used in studies of genetics, evolution, phylogeny, origin and taxonomy, as well as determining breeding strategies and the transfer of resistance genes through ploidy manipulation. However, chromosome identification is more of a challenge in many plant species with small chromosomes, such as potato (Solanum tuberosum L.). This is mainly due to difficulty in the preparation of good chromosome spreads, small chromosome size with similar morphology, and the low natural frequency of synchronously dividing cells in root-tip meristems. A high quality, well spread chromosome preparation is critical to enable routine work on potato ploidy manipulation, and physical gene localization by in situ hybridization for future advances in plant cytogenetics (Piñacker and Ferwerda, 1984; Sybenga, 1992; Visser et al., 1988; Wagenvoort et al., 1994).

Obtaining metaphase chromosomes from root tip cells is a common procedure in plant cytogenetics. Pretreatment of root tips with different reagents can dramatically affect the status of mitosis in root tips and influence the reliability of observations on the shape and size of chromosomes and the number of cells.

Materials and Methods

Potato materials used in this study included two accessions PI 275233 and PI 275236 from a wild diploid Mexican species Solanum pinnatisectum [Dunal (pmt, 2n = 2x = 24) and two breeding lines, CQ 123-25 and S 440 from the cultivated tetraploid potato S. tuberosum ssp. tuberosum (thr, 2n = 4x = 48). The plants from these four materials were maintained and propagated by tissue culture in MS medium (Murashige and Skoog, 1962). After 4 to 5 weeks of subcultures, the plants about 5 to 10 cm in height were transplanted into small pots (10 × 10 × 12 cm), which contained Cornell soilless mixture of 1 part of sphagnum peatmoss and 2 parts of horticultural grade vermiculite, and grown in a greenhouse under an 18-h light and 6-h dark photoperiod at 18 to 20°C. About 1 to 2 weeks after transplanting, the 1 to 2 cm long roots were collected from the plants by inverting the pots, and were dipped into 2 mL of pretreatment reagents contained in 1 cm diam test tubes. The root tips were sampled a total of three times at different dates spaced at 1-week intervals. For each sampling date, at least three root tips were collected from each line for every pretreatment. The results obtained from the three samplings on each line were combined to compare the different pretreatment methods for mitotic chromosome observation.

The four pretreatment methods used in this study included the following: T1, ice water for 24 h at 4°C; and T2, 10 ppm Ambush for 24 h at 4°C based on the treatment conditions described by Klein (1990). Ambush is a synthetic pyrethroid insecticide, a permethrin (Ambush, Pounce, Ectiban) [3-phenoxyphenyl methyl (±)-cis, trans-3- (2, 2-dichloroethenyl)-2, 2- dimethyl cyclopropene carboxylate] (ICI America, Inc, Wilmington, Del.); and T3, 2 mM 8-hydroxyquinolone for 4 h at 21°C according to the conditions described by Tjo and Levan (1950); and T4, CMMYT’s signal—a mixture containing 40 drops/100 mL of DMSO, 0.025% 8-hydroxyquinolone and 0.05% colchicine for 4 h at 21°C (Chen et al., 1998).

After pretreatment, the root tips were fixed in a 1:3 solution of glacial acetic acid: ethanol for 1 to 7 d at 4°C, followed by chromosome staining with 0.5% aceto-carmine for 10 to 20 min, and then root tip squashing in 45% acetic acid. Chromosome observation was conducted using an Olympus BX51 microscope with magnification ranging from 100X to 1000X. Comparison of the four pretreatment methods for chromosome observation was carried out by counting the total number of cells, the percentage of mitotic chromosomes at metaphase, and the percentage of countable mitotic cells with well spread chromosomes on a slide for each line. A mitotic index was determined using the following methods: 3 to 5 root tips per experiment for each line, and 250 cells per root tip were randomly analyzed. This procedure was repeated twice to determine the mean mitotic index (number of cells at metaphase × 100/total number of cells).

Based on the preliminary results, 1, 10, 100, 200, and 400 ppm of Ambush were evaluated to optimize the concentration of Ambush pretreatment.

Duncan’s multiple range procedure provided by the General Linear Model (GLM) program of SAS (SAS Institute Inc., Cary, N.C.) was used to compare the percentage of metaphase chromosomes and countable mitotic cells (P < 0.01).

Results

Table 1 summarizes the mitotic indices.
Table 1. Effect of different pretreatment methods on the mitotic index of root tips of genotypes from a wild diploid Mexican species, *S. pinnatisectum* (*pnt*, 2*n* = 2x = 24) and one cultivated tetraploid species *S. tuberosum* (*tbr*, 2*n* = 4x = 48).

| Solanum species | Treatment | Total Metaphase | Metaphase (%) | Countable | Countable (%) |
|-----------------|-----------|-----------------|---------------|-----------|---------------|
| *pnt*           | T1–ice water | 898 | 30 | 3.3 b<sup>z</sup> | 2 | 6.7 c |
|                 | T2–Ambush | 873 | 70 | 8.0 a | 28 | 40.0 a |
|                 | T3–8-Hydro | 855 | 20 | 2.3 b | 4 | 20.0 b |
|                 | T4–mixture | 883 | 7 | 0.8 c | 0 | 0.0 c |
| *tbr*           | T1–ice water | 860 | 32 | 3.7 b | 4 | 12.5 c |
|                 | T2–Ambush | 845 | 58 | 6.9 a | 32 | 55.2 a |
|                 | T3–8-Hydro | 905 | 13 | 1.4 c | 3 | 23.1 b |
|                 | T4–mixture | 899 | 13 | 1.4 c | 1 | 7.7 c |

<sup>z</sup>The results obtained from two wild diploid and two cultivated tetraploid potato lines were combined respectively.
<sup>y</sup>T1 = pretreatment with ice water for 24 h; T2 = 10 ppm Ambush for 24 h at 4 °C; T3 = 2 mM 8-hydroxyquinoline for 4 h at 21 °C; T4 = CIMMYT mixture solution for 4 h at 21 °C.
<sup>x</sup>Number of cells that were well spread and usable for chromosome counting.
<sup>z</sup>Means followed by different letters were significantly different at *P* < 0.01.

Based on the preliminary results, another experiment with the Ambush pretreatment at different concentrations (1, 10, 100, 200, and 400 ppm) was carried out to evaluate the effect of the concentration of Ambush on chromosome accumulation (Table 2). Concentrations of 1 to 100 ppm gave similarly high percentages of metaphase cells and countable cells, which were significantly superior to those of the 200 and 400 ppm treatments. Pretreatments with lower concentrations (1 and 10 ppm) of Ambush also provided the most distinguishable shapes and larger chromosomes in both of the *pnt* and *tbr* species (Fig. 1). Concentrations of more than 100 ppm resulted in overcondensed metaphase chromosomes, which might be desirable for chromosome counting, but were not suitable for banding or in situ hybridization studies where spatial resolution of the chromosomes is needed. It was noted that pretreatment with Ambush for 20 h at 4 °C resulted in longer chromosomes (Fig. 1B) than treatment for 24 h (Fig. 1A).

**Discussion**

The results presented showed that Ambush is an effective pretreatment reagent, producing better results for chromosome counts and cytogenetic analysis than the other three treatments. Using a concentration (1 to 10 ppm) of Ambush at a low temperature (about 4 °C) can significantly increase the mitotic index and produce the best separation of single chromosomes and well-spread metaphases (Table 1, Fig. 1). Using this technique, we could count 20 to 40 dividing cells for each slide preparation. This Ambush pretreatment method not only worked well on diploid species, but also on polyploid species, such as...
tetraploid potato with many chromosomes (Fig. 1C). Compared to traditional chromosome counting procedures, the present technique is simple, rapid and convenient for the detection and characterization of chromosomes in potato. Ambush pretreatment may be superior due to its disruption of karyokinetic spindles causing C-mitotic changes in plant mitotic cells (Klein, 1990). The inhibition of mitosis by Ambush, like many other alternative reagents including spindle poison herbicides (Dolezel et al., 1998), caused irregularities of the C-mitosis type, such as multinucleate cells and those containing restitution nuclei.

The present technique provides several advantages that make it suitable for cytogenetic study. The major advantages of Ambush in comparison with the other pretreatments, such as 8-hydroxyquinoline and cold water are as follows.

1) It provides higher quality of metaphase spreads. More than 50% of metaphases prepared using this method are well spread and usable for chromosome counting, whereas other pretreatments do not always show the desirable results.

2) The size and shape of chromosomes can be adjusted by using different concentrations and times of treatments. Usually, pretreatment with 1 to 10 ppm Ambush for a short time (about 20 h instead of 24 h) provides longer and distinguishable shaped chromosomes (Fig. 1). The longer chromosomes obtained using the present method (3.20 to 6.87 µm in pnt, data not shown), compared with those reported for small potato chromosomes (1 to 3.0 µm), would permit an accurate evaluation of chromosome morphology (Dong et al., 2000).

3) It is applicable to both diploid and polyploid species even though chromosome counting is always more difficult in polyploid species.

4) It can be applied to other plant species, especially those with many small chromosomes.

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