

Changes in Cellular Polyamine Contents and Activities of Their Biosynthetic Enzymes at Each Phase of the Cell Cycle in BY-2 Cells

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We analyzed changes in polyamine contents and the activities of biosynthetic enzymes during each phase of the cell cycle for a synchronized population of BY-2 cells. Based on our analysis of H³-thymidine incorporation flow cytometry, and the mitotic index, the M and G₂ phases seemed to occur at 8 h and from 2.5 to 8 h, respectively, after the release of aphidicolin. The respective activities of arginine decarboxylase (ADC), ornithine decarboxylase (ODC), and S-adenosyl methionine decarboxylase (SAMDC) at the beginning (7.4, 11.2, and 5.5 nmol mg⁻¹ protein h⁻¹) were increased to 22.6, 22.1, and 15.1 nmol mg⁻¹ protein h⁻¹. However, those increases do not coincide with the general change in polyamines reported from animal cells. In addition, the bi-phasic activation of polyamine biosynthetic enzymes, such as those found in the general animal model, was observed with ADC and ODC but not with SAMDC. These results suggest that the general animal model for explaining polyamine changes and SAMDC activation in the cell cycle cannot be applied to BY-2 cells. Further, our flow-cytometric analysis of cell populations may be a useful tool for evaluating the effects of polyamines on cell cycle progression in BY-2 cells.

Keywords: BY-2 cells, cell cycle, flow-cytometric analysis, *Nicotiana tabacum*, polyamines, S-adenosyl methionine decarboxylase

Cellular polycationic polyamines (spermidine, spermine, and their precursor, putrescine), which are ubiquitous in nature and absolutely required for eukaryotic cell growth, are synthesized in well-regulated biochemical pathways (Walden et al., 1997; Ackermann et al., 2003; Oredsson, 2003). The amino acids arginine and ornithine are converted to putrescine in a decarboxylation reaction catalyzed by arginine decarboxylase (ADC) and ornithine decarboxylase (ODC), respectively (Martin-Tanguy, 2001). Spermidine is synthesized in a reaction in which an aminopropyl group derived from decarboxylated S-adenosyl methionine (DcSAM) is coupled to putrescine by spermidine. A similar reaction is catalyzed by spermine synthase, converting from spermidine to spermine. DcSAM is generated by the action of S-adenosyl methionine decarboxylase (SAMDC), which is the rate-limiting step in the polyamine biosynthetic pathway (Anderson et al., 1998; Martin-Tanguy, 2001; Ackermann et al., 2003).

Polyamines promote plant growth and development by activating the synthesis of nucleic acids and proteins (Amarantos et al., 2002; van Dam et al., 2002). They have been implicated in such processes as cell proliferation, differentiation, flowering, and

senescence (Walden et al., 1997; Martin-Tanguy, 2001; Jang et al., 2002; Ahn and Jin, 2004). Cell proliferation ensures the production of two new healthy daughter cells at the end of a cell cycle (Oredsson, 2003). To understand the roles of polyamines in that cycle, researchers use either the inhibitors of various polyamine biosynthetic enzymes, e.g., difluoromethyl arginine (DFMA), difluoromethyl ornithine (DFMO), and methyl-glyoxal-bis-guanylhydrazone (MGBG), or mutants and transgenic cells that lack polyamine biosynthetic genes, such as ADC, ODC, or SAMDC. For example, cells of *Schizosaccharomyces pombe*, which cannot synthesize spermidine or spermine because of a deletion-insertion in the gene coding for SAMDC, show an overall delay in cell cycle progression and an accumulation of cells in the G₁ phase (Chattopadhyay et al., 2002). In addition, polyamine starvation via inhibition of biosynthetic enzymes prolongs the S and G₂ phases in polyamine-dependent Chinese hamster ovary cells (Anehus et al., 1984). ODC and SAMDC activities, as well as polyamine contents, have been investigated during the cell cycle in populations synchronized by various methods. For animal systems, the

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Abbreviations: ADC, arginine decarboxylase; DcSAM, decarboxylated S-adenosyl methionine; ODC, ornithine decarboxylase; SAM, S-adenosyl methionine; SAMDC, S-adenosyl methionine decarboxylase.

general conclusions are that ODC and SAMDC are activated in a bi-phasic manner, with a first activation phase in late G₁ close to the G₁/S transition, and then a second activation phase in conjunction with S/G₂ transition. Moreover, the doubling of putrescine, spermidine and spermine contents mainly occurs during the late S phase, S/G₂ transition, and the G₁ and S phases (Oredsson, 2003).

The most easily synchronizable plant system (Ehsan et al., 1999) is from a tobacco (*Nicotiana tabacum* L. cv. Bright Yellow-2) cell line (Nagata et al., 1992; Nagata and Kumagai, 1999). Those cells have been used successfully to study the effects of stress on cell cycle progression (Reichheld et al., 1999; Świąte et al., 2002). For example, the expression of cyclin-dependent kinases (CDKs) in the plant cell cycle has been examined in synchronized BY-2 cells (Setiady et al., 1996; Reichheld et al., 1999; Sorrell et al., 2001; Świąte et al., 2002). Likewise, flow-cytometrical analysis also has established a cell cycle for BY-2 cells (Świąte et al., 2002).

Plant cells store polyamines in their vacuoles, but ADC also has a minor role in the synthesis of putrescine in animal cells (Martin-Tanguy, 2001). Antibodies against oat ADC have revealed that this enzyme is present in the thylakoid membranes of chloroplasts (Borrell et al., 1996). SAMDC and spermidine synthase are localized to the cytoplasm, whereas ODC is found in the nucleus (Martin-Tanguy, 2001). Additional functions for plant polyamines may include the maintenance of photosynthetic activity and the prevention of osmotic stress-induced senescence (Borrell et al., 1996; Martin-Tanguy, 2001), both of which not occurring in animal cells. These reports support the possibility that metabolism differs between plants and animals for polyamine biosynthesis or storage. No reports have been published about how plant polyamines and their biosynthetic enzymes can regulate each phase of the cell cycle. Therefore, the objective of our current work was to clarify the changes in polyamine contents and the activation of their biosynthetic enzymes during cell cycle progression in BY-2 cells. Here, we present data on the effect of exogenous polyamine on that progression, as well as changes in ADC, ODC, and SAMDC activities and polyamine contents at various cycle checkpoints.

MATERIALS AND METHODS

Chemical Reagents and Protein Determinations

Standard chemicals were purchased commercially.

Protein contents were estimated according to the method of Bradford (1976), as described by Wi and Park (2002).

Cell Culture, Synchronization, and Flow-cytometrical Analysis

Tobacco BY-2 suspension cells were used. Cell maintenance and synchronization with aphidicolin at the G₁/S phase were achieved as described by Nagata et al. (1992). The flow-cytometric analysis was executed according to Świąte et al. (2002). Suspension cells were washed with 0.66 M sorbitol in an MS (Murashige and Skoog) medium, and then incubated with 0.1% (w/v) pectolyase and 2% (w/v) cellulase for 1 h at 37°C. After isolation via centrifugation, DNA from the released nuclei was stained with a solution containing propidium iodide, and analyzed on an FACS analytical flow cytometer (Coulter, USA). To investigate the effects of exogenous polyamines on the cell cycle of BY-2 cells, subcultures were treated with 1 mM of putrescine, spermidine, and spermine for 12 h. Fractionation of each phase followed the flow-cytometric data, according to the method of Menges and Murray (2002).

Determination of Polyamine Contents and Activities of Biosynthetic Enzymes

Polyamines were analyzed as described by Wi and Park (2002). The supernatant of leaf extracts (0.2 mL) was mixed with saturated sodium carbonate (0.2 mL) and dimethylaminonaphthalene-1-sulfonyl chloride (0.4 mL), and then incubated at 60°C for 1.5 h. Afterward, the dansylated products were extracted with benzene. Aliquots were separated by thin layer chromatography in 25:2 (v:v) chloroform:triethylamine. The separated polyamines were quantified on an RF-1501 (Shimadzu, Japan) spectrofluorophotometer (wavelengths of excitation/emission: 350/495 nm). Enzyme activities were determined according to the method of Park and Lee (1994). Briefly, frozen powder that had been ground in liquid nitrogen was re-suspended in a buffer containing 25 mM Tris-Cl (pH 7.6), 1 mM EDTA, and 15 mM β-mercaptoethanol, using a ratio of 2:3 (w/v) for tissue to buffer. The supernatant fraction served as the enzyme source for measuring the activities of SAMDC, ODC, and ADC. A 300 μL aliquot of the extract, containing 0.1 μL of [carboxyl-¹⁴C] SAM, 0.05 or 0.1 μCi of L-[1-¹⁴C] Orn, and 0.1 μCi of DL-[1-¹⁴C] Arg, was added to a series of flasks with center wells into which were placed filter papers

with 2 N KOH. After incubation at 37°C for 60 min, the reaction was stopped with 5% TCA. The filter paper that had adsorbed the $^{14}\text{CO}_2$ was evaluated for radioactivity in a liquid-scintillation counter.

RESULTS AND DISCUSSION

Cyclin A and B are Specifically Expressed in the S and G₂/M Phases, Respectively, of the BY-2 Cell Cycle

Phase progression in the cell cycle of BY-2 cells was analyzed by flow cytometry (Fig. 1A); the results produced were similar to those previously reported with *Arabidopsis* (Porceddu et al., 2001) and other BY-2 cells (Jang et al., 2005). Quantification of ratios for individual cycle phases was based on data from the flow-cytometrical analysis (Fig. 1B), and showed that 69.8% of the cells were in the G₂ phase at 3 h after the release of aphidicolin. The respective percentages for cells in the G₁ and G₂ phases versus total cells dramatically changed from 1.5% and 69.7% at hour 6 to 59.8% and 27.2% at hour 9. Our results for ^3H -thymidine incorporation and the change in mitotic index (Fig. 2A) subsequently confirmed those obtained from the flow cytometry (Fig. 1B). Therefore, we can

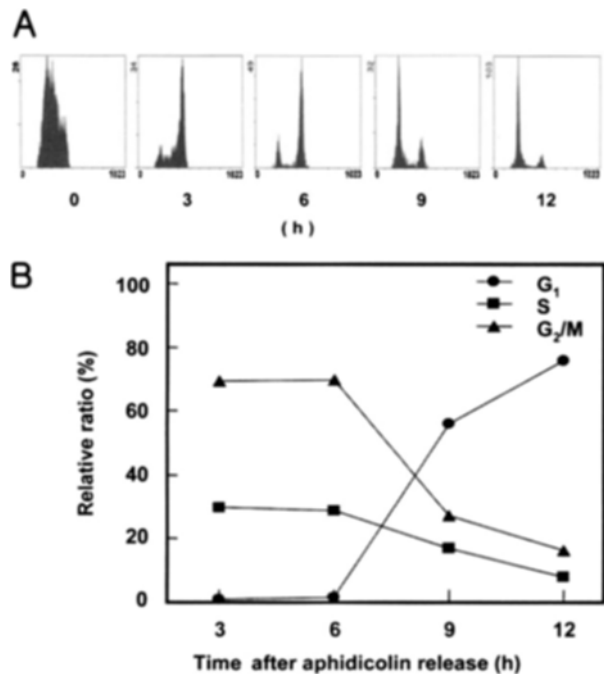


Figure 1. Changes in population of BY-2 cells after the release of aphidicolin. **A**, Flow-cytometric analysis. **B**, Relative ratio of cell population for each incubation period.

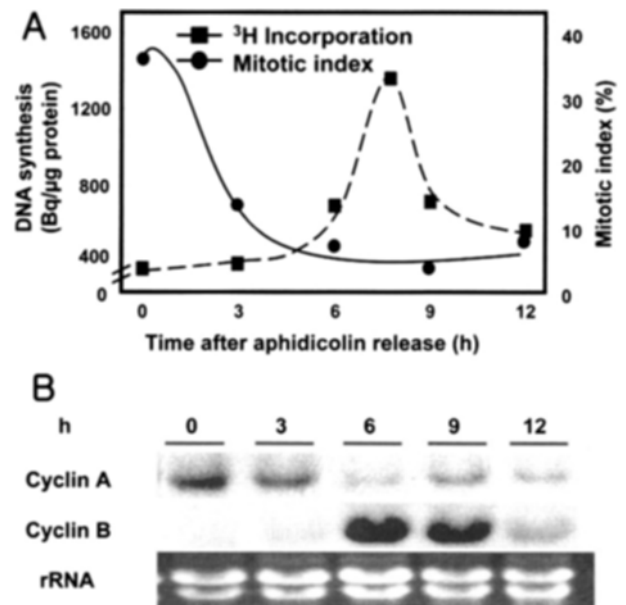


Figure 2. Changes in indicators that separate phases in cell cycle after the release of aphidicolin. **A**, ^3H -thymidine incorporation and mitotic index. **B**, Expression of cyclins A and B.

conclude that the G₂ and M phases in BY-2 cells was at hour 3 and 8.

We also monitored changes in the expression of cyclin A (GenBank accession No. AF518250) and B (GenBank accession No. AF518251). Our data for the expression of cyclin A in the S phase (Fig. 2B) supports a previous result that A-type cyclin plays an essential function in DNA replication during that phase (Reichheld et al., 1999). In addition, the expression pattern observed here for cyclin B in the G₂/M phase coincides with the previous report (Sorrell et al. 2001).

Exogenous Polyamines Stimulate Accumulation of Cyclin A and B in BY-2 Cells

Based on our current data, the question arises as to whether flow cytometry could adequately analyze the changes in each cell cycle phase for BY-2 cells. Although polyamine starvation can prolong or arrest specific phases (Anehus et al., 1984; Chattopadhyay et al., 2002), treatment with polyamine can stimulate the progression of the cell cycle (Pohjanpelto et al., 1994). Therefore, it is possible that polyamine can control that progression. We conducted a series of experiments that assessed the cell population at each phase in the cycle, and compared the expression levels of cyclin A and B in BY-2 cells treated with polyamines. Exogenous applications did indeed

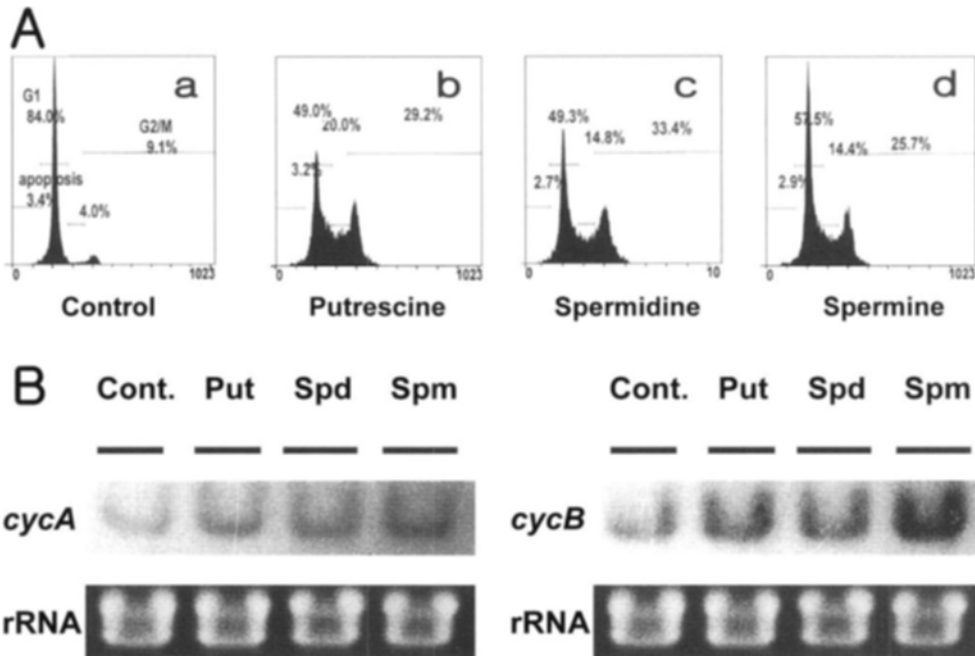


Figure 3. Effects of exogenous polyamines during phases of cell cycle. **A**, Changes in Polyamine in G₁, S, and G₂ phases from 4-day-old BY-2 cells. **B**, Effect of exogenous polyamines on expression of cyclins A and B in 4-day-old BY-2 cells.

induce the relative accumulation of cells during the S and G₂/M phases (20% of S and 28% of G₂ for putrescine, 14% of S and 33.4% of G₂ for spermidine, and 14% of S and 25.7% of G₂ for spermine) compared with the corresponding control (4.2% of S and 9.1% of G₂) (Fig. 3A). We also performed northern analysis of cyclin A and cyclin B genes, which were specifically expressed in the S and G₂/M phase, respectively, to confirm this accumulation while BY-2 cells were being incubated with exogenous polyamines. The ratios for cell populations determined by flow-cytometric phase separation coincided with the accumulation of both genes (Fig. 3B). These results support that measurements of cell populations via flow-cytometric phase separation can adequately describe changes in each phase of the cell cycle for BY-2 cells. In addition, our findings may extend to the control of gene expression related with cell cycle progression by exogenous polyamines.

Polyamine Content Depends on the Cell Cycle

Using the results shown in Figure 1 and 2, we constructed a scheme to illustrate phase separation during the cell cycle (Fig. 4A). Polyamines are essential for both the synthesis of DNA in the S phase and the stabilization of the DNA structure (Fillingame et al., 1975; van Dam et al., 2002), as well as for cell

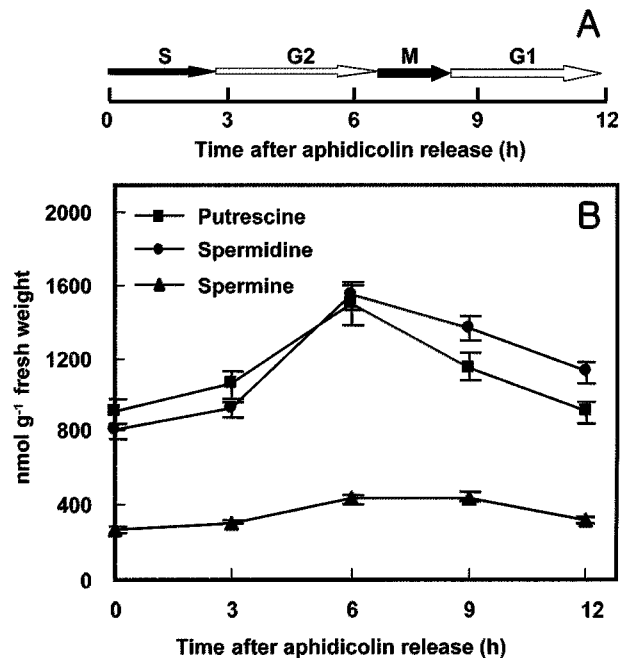


Figure 4. Comparison between phase separation scheme of aphidicolin-based synchronization **A** and changes in polyamines after releasing aphidicolin **B**.

growth and proliferation (Martin-Tanguy, 2001). These reports and our results in Figure 3 confirmed that cellular polyamines play an important role in the pro-

gression of the cell cycle. Because polyamine contents had not previously been analyzed in each phase, we did so here, and found that polyamine increased in the G₂ phase (Fig. 4B), which concurred with our schematic depiction of phase separations based on the time that elapsed after aphidicolin was released (Fig. 4A). Thus, about twice as much putrescine, spermidine and spermine were accumulated in the S (or late G₁) vs. the G₂ phases, whereas the ratio of (spermidine + spermine)/putrescine seemed to remain constant at each phase. In contrast, the doubling phase in animal systems differs, i.e., late S phase for putrescine, S/G₂ transition for spermidine, and G₁ and S phases for spermine (Oredsson, 2003). Therefore, our data support the conclusion that regulation of polyamine contents in BY-2 cells cannot and should not be explained by the general animal model (Oredsson, 2003).

Activations of Polyamine Biosynthetic Enzymes Depend on the Cell Cycle and SAMDC

To test whether changes in cellular polyamine were a direct result of enzymatic activation, we analyzed activities of key enzymes for polyamine biosynthesis. At the starting point, ADC, ODC, and SAMDC activities were 7.4, 11.2, and 5.5 nmol mg⁻¹ protein h⁻¹; at their peaks, these levels were 22.6, 22.1, and 15.1 nmol mg⁻¹ protein h⁻¹, respectively (Fig. 5A). However, compared with the phase period after the release of aphidicolin, peaks in ADC and ODC activities were localized to the G₂ phase (Fig. 5A). This indicates that the observed increase in putrescine at the G₂ phase may have directly resulted from the rise in ADC and ODC activities. In contrast, the doubling in levels of spermidine and spermine was not observed at the peak of SAMDC activity, which occurred 3 h earlier than for the others (Fig. 4B, 5A). Moreover, when the changes in ADC, ODC, and SAMDC activities were re-drawn from G₁ to M, ADC and ODC were activated in a bi-phasic manner during cell cycle progression. The first and second activations of ADC and ODC were at G₁/S and G₂, respectively, which coincides with the activation of ODC reported in animals (Oredsson, 2003). However, SAMDC activity in our BY-2 cells peaked, without marked fluctuation, during the S/G₂ transition. This suggests that cell-cycle changes in SAMDC activity may differ from those reported for the general animal model (Oredsson, 2003).

As analyzed by flow cytometry, the coincidence noted between the expression of cyclin (A and B)

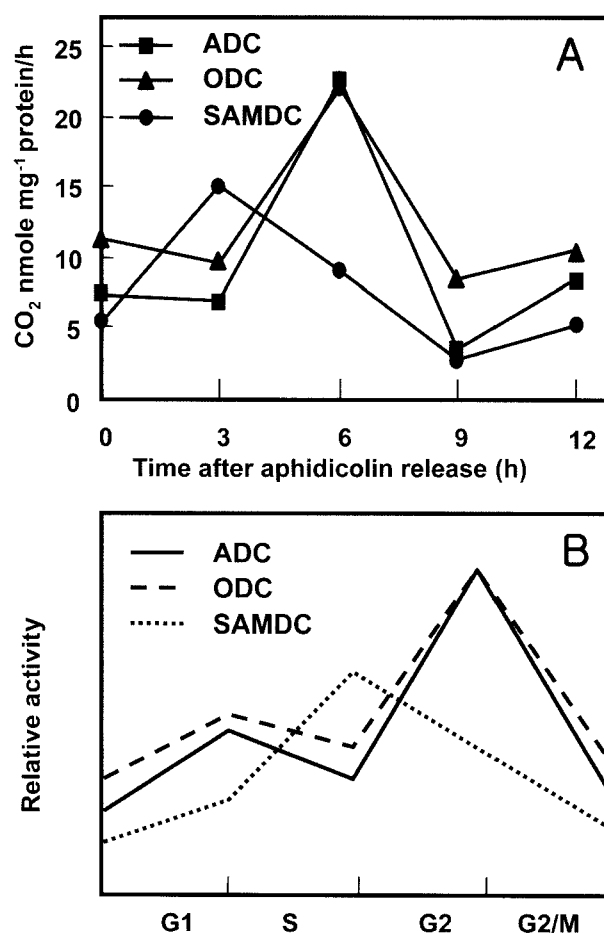


Figure 5. Fluctuations in levels of polyamine biosynthetic enzymes after the release of aphidicolin. **A**, Activities of ADC, ODC, and SAMDC. **B**, Schematic presentation of changes in ADC, ODC, and SAMDC activities with orderly progression of cell cycle. (Phase separation from G₁ to M was based on data in Figure 4A; relative activity is ratio of highest activity among ADC, ODC, and SAMDC).

with/without polyamine and the ratio of cell populations in each phase raises the possibility of using synchronized BY-2 cells to identify the role of polyamine in controlling the cell cycle in plants. In addition, our present data demonstrate that the general animal model (Oredsson, 2003) should not apply to BY-2 cells for explaining polyamine changes and SAMDC activation in the cell cycle.

Based on our knowledge, SAMDC activity in carnation may be controlled by transcriptional regulation (Mad Arif et al., 1994; Lee et al., 1997; Park et al., 2001; Kim et al., 2004) or translational regulation with uORF (Raney et al., 2002). This SAMDC activity is also highly regulated by a variety of physiological, hormonal, and environmental stimuli (Lee et al., 1996; Walden et al., 1997; Martin-Tanguy, 2001;

Marco and Carrasco, 2002). Therefore, our current work suggests three questions. First, can the mechanics for cyclin-dependent kinase regulation of the cell cycle (Menges and Murray, 2002) also be used to explain the changes in SAMDC activity during the cell cycle in plants? Second, is the increase in SAMDC activity related to translational inhibition by uORF? Finally, how do changes in endogenous polyamine, due to the over-expression of SAMDC, affect cell cycle progression in BY-2 cells? Further research should involve the use of SAMDC gene-transformed BY-2 cells to determine the role of SAMDC in the cell cycle.

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