

DIFFERENTIAL CELL CYCLE PERTURBATION BY TRANSMETHYLATION INHIBITORS

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Abstract—Cell cycle distribution of HL-60 cells was studied by flow cytometry after incubation with the transmethylation inhibitors 3-deaza-(\pm)-aristeromycin (c^3 Ari) and 3-deazaadenosine (c^3 Ado). Cells were incubated with the drugs (25 μ M) for two cell doublings in control cells (36 hr). The presence of c^3 Ari caused a dose-dependent, reversible G_2 +M arrest, whereas c^3 Ado-treated cells accumulated in G_0 / G_1 . The G_2 +M arrest was also found in NIH/3T3 cells incubated for 36 hr with 25 μ M c^3 Ari, but not in U937 and K562 cells. Possible mechanisms for the described effects of c^3 Ari are discussed from the perspective that inhibition of *S*-adenosyl homocysteine hydrolase, and subsequent inhibition of transmethylation reactions, at present is the only known site of action of c^3 Ari.

The adenosine analogues c^3 Ado† and c^3 Ari have been extensively studied as inhibitors of AdoHcyase (E.C.33.22.1) and transmethylation reactions [1–3]. The diverse biological effects of c^3 Ado have been associated with interactions with a number of primary cellular targets, while the effects of c^3 Ari have not been ascribed to interaction with any cellular component apart from AdoHcyase [3]. Thus, c^3 Ado and c^3 Ari emerge as examples of non-selective and selective transmethylation inhibitors, respectively.

We have previously investigated the capacity of c^3 Ado and c^3 Ari as inducers of differentiation of leukemia cell lines *in vitro* [4, 5] and the effects of c^3 Ado on viability and growth of human cells potentially involved in differentiation induction *in vivo* in man [6]. Herein, we present differential effects of c^3 Ado and c^3 Ari on cell cycle.

MATERIALS AND METHODS

Cells. Four different cell lines were tested: HL-60 (a human promyelocytic leukemia cell line), U937 (a monocytelike cell line derived from a patient with histiocytic lymphoma), K562 (a human myeloid leukemia-derived cell line) and NIH/3T3 (a murine fibroblast cell line). Cells were generous gifts from M.S. Hersfield, Duke University, U.S.A. (HL-60), S.O. Kolset, University of Tromsø (U937, K562) and I.W. Nilsen, University of Tromsø (NIH/3T3). Cells were grown in RPMI 1640 (Gibco Ltd, Paisley, U.K.) supplemented with 10% heat inactivated horse serum (HL-60), fetal calf serum (U937) or newborn calf serum (K562). NIH/3T3 cells were grown in Dulbecco's (Gibco) medium supplemented with 10% heat inactivated fetal calf serum. All cells were grown in a fully humidified atmosphere of 5% CO₂. Cell counts were determined in duplicate by a

hemocytometer, and cell viability was assessed by trypan blue exclusion.

Experiments. Suspensions of cells were adjusted to concentrations of early logarithmic growth. The following day, cell suspensions were transferred to 50 mL culture flasks, and drug or medium added to a final volume of 10 mL. After each incubation interval, one set of flasks was harvested and two 100 μ L samples were obtained for determination of cell concentration. The remaining cells were used for DNA-analysis.

NIH/3T3 cells were harvested by trypsinization and resuspended in 10 mL of medium followed by the same procedure as for the other cultures.

HL-60 cells exposed to 25 μ M c^3 Ari were washed once in drug-free medium before adjustment of cell concentration and reincubation in drug-free medium.

DNA-analysis. Cells were centrifuged for 5 min at 200 g, washed in phosphate buffered saline (Dulbecco), resuspended in saline, fixed in 70% ethanol and kept refrigerated until analysis. Twenty-four hours before analysis, cells were heated at 45° for 30 min [7], centrifuged, and stained with 50 μ g/mL Mithramycin (Pfizer, NY, U.S.A.) in 25% ethanol-solution containing 15 mM MgCl₂. Cells were subsequently kept refrigerated until analysed on an Argus Flowcytometer (Skatron, Lierbyen, Norway) equipped with an Ahrens FDAS data system (O. Ahrens, Bargteheide, F.R.G.). The number of cells in different cell cycle phases were calculated from the DNA-histograms by means of the peak reflect method [8]. The results are given as percentage of total histogram cell count. Calculations for statistical significance by Student's *t*-test were done on a microcomputer with the Microstat (Eco-soft, INC., U.S.A.) program.

RESULTS

After more than 36 hr incubation, 25 μ M c^3 Ado decreased the number of viable HL-60 cells compared to controls (Fig. 1). In cells incubated with

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† Abbreviations: c^3 Ado, 3-deazaadenosine; c^3 Ari, 3-deaza-(\pm)-aristeromycin; AdoHcyase, *S*-adenosyl homocysteine hydrolase; Adomet, *S*-adenosyl homocysteine.

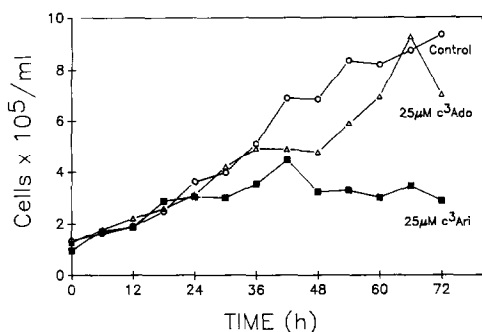


Fig. 1. Growth of HL-60 cells in the presence of 3-deazaadenosine and 3-deaza-(±)-aristeromycin.

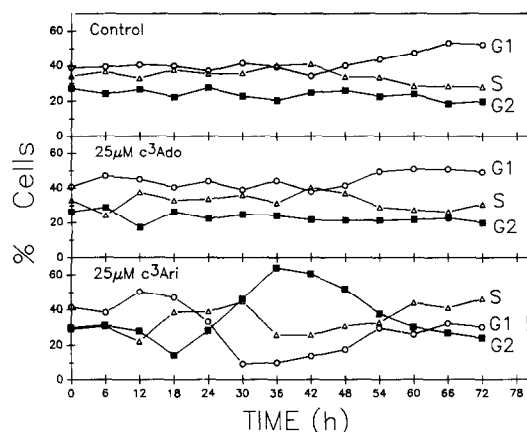


Fig. 2. Flow cytometric analysis of cell cycle distribution of HL-60 cells during exposure to 3-deazaadenosine and 3-deaza-(±)-aristeromycin.

25 μM c³Ari, growth was inhibited after 24 hr incubation.

Cells incubated with 25 μM c³Ari showed a different pattern of cell cycle distribution as compared to control cells, and c³Ado-exposed cells incubated for more than 18 hr (Fig. 2). The most striking effect of c³Ari was a rise in the percentage of cells in G₂+M, which reached a maximum of more than 50% of total after 36 hr (Fig. 3) and decreased to control values after 72 hr of incubation. Notably, the effect was also reflected in an increase in the absolute number of cells in G₂+M. c³Ari induced a decrease in the percentage of cells in G₀/G₁ after 18 hr with a trough value of about 10% at 30 hr, followed by a slow subsequent increase. The percentage of cells in S increased slightly between 18 and 30 hr of incubation, approaching control levels at 54 hr, with a slow increase from 36 hr onwards. The endpoint distribution represents a shift in percentage of cells in S and G₀/G₁ compared to control cells.

Table 1 summarizes the results from five experiments with HL-60 cells incubated for 36 hr. Cells incubated with 25 μM c³Ari demonstrates a G₂+M arrest while the most striking effect of 25 μM c³Ado after 36 hr was an increase in the percentage of cells in G₀/G₁.

After 36 hr incubation with c³Ari, effects were seen on cell cycle distribution at concentration of 5 μM, with a dose-dependent redistribution reaching a plateau at 40 μM (Fig. 4).

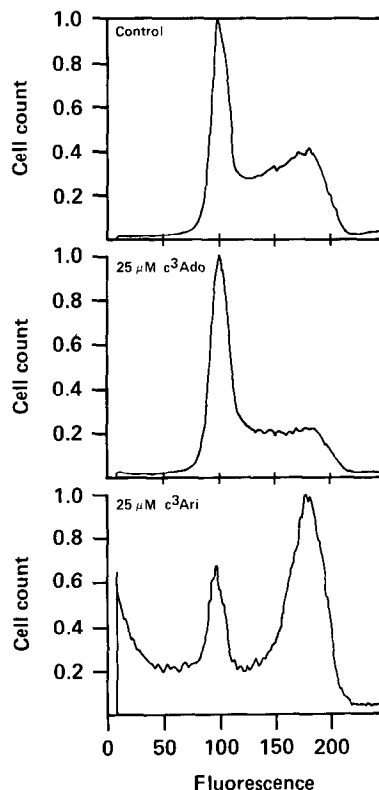


Fig. 3. DNA histograms of HL-60 cells after 36 hr exposure to 3-deazaadenosine and 3-deaza-(±)-aristeromycin. Cells were fixed in 70% ethanol and stained with mithramycin.

HL-60 cells exposed to 25 μM c³Ari for 36 hr regained control distribution after approximately 18 hr incubation in drug-free medium (Fig. 5). Compared to control cells, growth was slightly reduced in the pre-exposed cells after 24 hr of drug-free incubation (Data not shown).

A limited number of cell lines were investigated to see if the cell cycle effects of 25 μM c³Ari for 36 hr in HL-60 cells could be reproduced using other target cells. Only NIH/3T3 cells showed a significant increase in the percentage of cells in G₂+M and a reduction in G₀/G₁ (Table 2).

DISCUSSION

The data presented herein are the first demonstration of cell cycle effects of transmethylation inhibitors. The effects of c³Ari are particularly interesting because the compound appears to be a selective inhibitor of AdoHcyase. Most known inducers of leukemia cell differentiation cause cell arrest in G₀/G₁. Dimethylsulphoxide and phorbol esters induce HL-60 arrest in G₀/G₁ after one doubling [9, 10]. Isotretinoin treatment causes a similar arrest after two divisions [9]; while sodium butyrate blocks cells in late G₁ [11]. Among agents known to induce differentiation *in vitro*, the G₂+M arrest seems to be confined to inhibitors of mitosis [12].

The present data show that a proposed selective transmethylation inhibitor, c³Ari, also reversibly arrests HL-60 cells in G₂+M. At the dose schedule

Table 1. Cell cycle distribution of HL-60 cells incubated with 25 μ M 3-deazaadenosine and 3-deaza-(\pm)-aristeromycin for 36 hr

	% Cells in G ₀ /G ₁	% Cells in S	% Cells in G ₂ +M	Growth as % of control
Control	34.3 \pm 1.3	36.0 \pm 1.2	29.7 \pm 0.4	—
25 μ M c ³ Ado	47.5 \pm 1.9*	32.3 \pm 1.3*	20.2 \pm 0.7*	90.9 \pm 11.2
25 μ M c ³ Ari	21.9 \pm 1.5*	26.6 \pm 1.8*	51.6 \pm 0.8*	60.7 \pm 5.2*

Results are given as mean \pm SD from five experiments.

* $P < 0.05$ by Student's t -test.

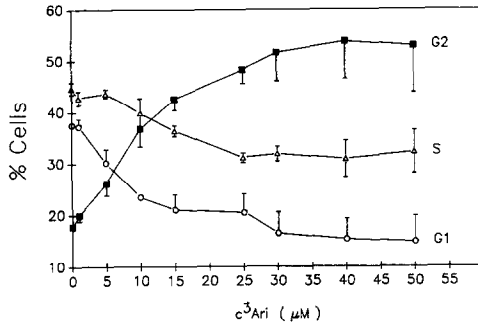


Fig. 4. Dose-response relationship for HL-60 cell cycle distribution after 36 hr exposure to 3-deaza-(\pm)-aristeromycin.

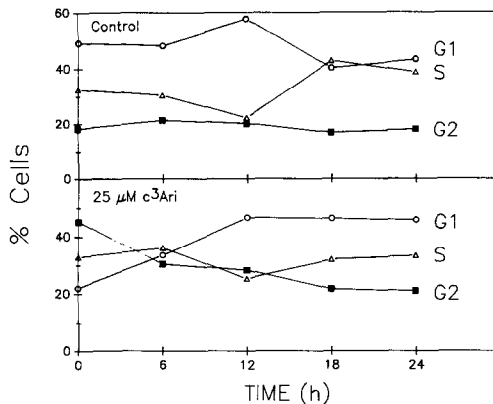


Fig. 5. Cell cycle distribution after washout of 3-deaza-(\pm)-aristeromycin. HL-60 cells were incubated with either medium alone (control) or with 3-deaza-(\pm)-aristeromycin for 36 hr, which is time zero on the figure. At this point medium was removed and cells reincubated in drug-free medium.

employed, this effect of c³Ari could not be demonstrated in two other human hemopoietic cell lines. However, this effect of c³Ari is not confined to human hemopoietic cell lines undergoing differ-

entiation, as demonstrated by the G₂+M arrest in the murin fibroblast cell line NIH/3T3.

The G₂+M arrest in HL-60 was reversible both after continuous exposure to c³Ari and after washout of c³Ari. We are presently investigating how these effects are interrelated.

Differential effects on cell cycle can be included on the list of differences between the AdoHcyase inhibitors c³Ado and c³Ari. Only c³Ado has been shown to affect cyclic AMP levels [13, 14] and microfilaments [15]. While both c³Ado [16] and c³Ari [17] are phosphorylated to the monophosphate in intact cells, only c³Ado has been detected as the triphosphate [16, 17]. In addition, only c³Ado commits HL-60 cells to differentiation after a short time (6 hr) of exposure [18], while continuous presence of c³Ari is necessary to induce differentiation [5], and only c³Ado modulates cellular glutathione levels [19]. The observation that cell cycle effects of c³Ari are exerted at concentrations known to effectively inhibit HL-60 AdoHcyase [6], is consistent with the concept of c³Ari as a selective transmethylation inhibitor. The G₂+M arrest is reversible upon removal of c³Ari consistent with the c³Ari effects reported for growth inhibition [6] and HL-60 cell differentiation [18]. Based on the present knowledge, it may seem reasonable to assume that c³Ari induces G₂+M arrest by a mechanism related to inhibition of AdoHcyase.

Three possible relations have been partly explored. Firstly, c³Ari could block increased activity of AdoHcyase needed for traverse through G₂+M. Although the activity of the enzyme in mouse mastocytoma P-815 cells and its c-AMP binding capacity were at a maximum in G₂+M [20], the activity of AdoHcyase showed no cell cycle dependency in untreated HL-60 cells [21].

Secondly, enzyme activities and cellular metabolites of general importance for transmethylation reactions could show variation through the cell cycle, and cells could be particularly vulnerable to perturbation of AdoHcyase inhibition in G₂+M. The

Table 2. Cell cycle distribution and growth of cells incubated with 25 μ M 3-deaza-(\pm)-aristeromycin for 36 hr

	% Cells in G ₀ /G ₁	% Cells in S	% Cells in G ₂ +M	Growth as % of control
3T3 Control	32.9 \pm 6.2	39.6 \pm 5.9	27.5 \pm 0.7	—
3T3 25 μ M c ³ Ari	14.3 \pm 0.5*	36.4 \pm 4.3	49.3 \pm 4.1*	53.3 \pm 10.8*
K562 Control	37.0 \pm 5.4	36.8 \pm 5.7	26.1 \pm 6.1	—
K562 25 μ M c ³ Ari	34.5 \pm 6.3	39.9 \pm 5.9	25.6 \pm 8.2	84.5 \pm 17.8
U937 Control	38.3 \pm 5.4	37.6 \pm 3.5	24.1 \pm 6.1	—
U937 25 μ M c ³ Ari	42.5 \pm 3.4	36.8 \pm 1.1	20.7 \pm 2.3	58.0 \pm 22.8*

Results are given as mean \pm SD from three experiments.

* $P < 0.05$ by Student's t -test.

activity of *S*-adenosyl methionine synthase did not, however, show cell cycle dependency in HL-60 cells, nor did AdoMet and AdoHcy concentrations, or the AdoMet/AdoHcy ratio [22].

Thirdly, *c*³Ari could inhibit specific methylation reactions, like DNA methylation. Although *c*³Ari did not change overall DNA methylation [23], a comprehensive exploration of the concept that *c*³Ari may act on cell cycle distribution through transmethylation inhibition, should include the possibility that *c*³Ari could alter sequence specific promoter methylation of genes [24], e.g. inhibitory genes involved at a restriction point in late G₂ [25].

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