

Influence of LOX/COX inhibitors on cell differentiation induced by all-*trans* retinoic acid in neuroblastoma cell lines

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Abstract. We investigated the possible modulation by LOX/COX inhibitors of all-*trans* retinoic acid (ATRA)-induced cell differentiation in two established neuroblastoma cell lines, SH-SY5Y and SK-N-BE(2). Caffeic acid, as an inhibitor of 5-lipoxygenase, and celecoxib, as an inhibitor of cyclooxygenase-2, were chosen for this study. The effects of the combined treatment with ATRA and LOX/COX inhibitors on neuroblastoma cells were studied using cell morphology assessment, detection of differentiation markers by immunoblotting, measurement of proliferation activity, and cell cycle analysis and apoptosis detection by flow cytometry. The results clearly demonstrated the potential of caffeic acid to enhance ATRA-induced cell differentiation, especially in the SK-N-BE(2) cell line, whereas application of celecoxib alone or with ATRA led predominantly to cytotoxic effects in both cell lines. Moreover, the higher sensitivity of the SK-N-BE(2) cell line to combined treatment with ATRA and LOX/COX inhibitors suggests that cancer stem cells are a main target for this therapeutic approach. Nevertheless, further detailed study of the phenomenon of enhanced cell differentiation by expression profiling is needed.

Introduction

Neuroblastoma is the most common extracranial malignant solid tumor in children, accounting for 6-8% of all childhood cancers and more than 15% of pediatric cancer deaths (1). Neuroblastomas are neuroectodermal tumors arising from elements of the neural crest. The clinical behavior of neuroblastoma is heterogeneous; the likelihood of survival depends

on the age at diagnosis, extent of disease and tumor biology. Three different patterns are noted: life-threatening progression with fatal outcome, maturation into ganglioneuroblastoma or ganglioneuroma (even metastatic neuroblastomas have been shown to spontaneously mature to benign ganglioneuromas), or spontaneous regression (2-4).

Approximately half of the patients are diagnosed with high-risk disease, with overall survival rates less than 40% despite intensive multimodal therapy. In general, the clinical outcome of children with high-risk or relapsed neuroblastoma is poor despite the introduction of dose-intensified and high-dose chemotherapy (1,5). The toxicity of current treatment protocols remains significant, and there is little space to further intensify therapy. Alternative treatment strategies are thus needed in order to improve survival as well as to diminish late effects of therapy. Besides immunotherapy, which offers a much more specific and less toxic treatment than conventional therapies (6), treatment strategies that target multiple levels including tumor cells, stromal cells and angiogenesis may be of clinical benefit (7-9).

Induced differentiation of transformed cells into mature phenotypes represents a promising strategy in recent anti-tumor therapy (10-13). Among various inductors of differentiation, retinoids are the most frequently investigated. These derivatives of vitamin A participate in morphogenesis and differentiation processes during mammalian development. In particular, retinoic acid (RA) plays an important role in the proliferation and differentiation of many cell types and can reverse malignant growth both *in vitro* and *in vivo* (14-16).

It is evident that the biological effects of retinoic acid are mediated principally through retinoic acid receptors (RARs) and retinoid X receptors (RXRs). These receptors represent two subgroups of steroid and thyroid hormone receptor families that form receptor dimers, which then bind to specific response elements and regulate gene transcription (17). The role of retinoids in transcription activation machinery is particularly influenced by retinoic acid stereoisomers, i.e., all-*trans* retinoic acid (ATRA), 13-*cis* retinoic acid (13-*cis* RA) and 9-*cis* retinoic acid (9-*cis* RA), which are synthesized from retinaldehyde precursors. Among these, ATRA is the most abundant and thermodynamically stable form (18).

Studies comparing the activity of 13-*cis* RA and ATRA in neuroblastoma cell lines have demonstrated similar potency

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in terms of cell differentiation, growth arrest and regulation of gene expression (19). Furthermore, several lines of evidence suggest that 13-*cis* RA is likely to isomerize *in vivo* to ATRA and also to 9-*cis* RA; these are now believed to be the main biologically active forms of RA, although little is known about the mechanisms of RA isomerization *in vivo* (18,20,21).

In the past few years, attention has also been paid to the possibility of combined induction of differentiation as well as to the modulation of differentiation by other compounds. This therapeutic approach was reported particularly for leukemia treatment; however, this strategy was also demonstrated to be effective *in vitro* with various types of cell lines derived from pediatric solid tumors. It has been reported that ATRA-induced differentiation in the HL-60 leukemia cell line can be further enhanced by combined application with bile acids (22,23). Retinoid-induced differentiation of neuroblastoma cells may be influenced by inhibitors of enzymes that participate in the intracellular degradation pathway of retinoids, the arachidonic acid metabolic pathway, especially inhibitors of lipoxygenases (LOX) and cyclooxygenases (COX) (24,25). Our study is therefore focused on the influence of arachidonic acid metabolic pathway inhibitors on ATRA-induced differentiation in neuroblastoma cell lines.

The aim of this study was to investigate a possible effect of arachidonic acid metabolic pathway inhibitors on the differentiation of SH-SY5Y and SK-N-BE(2) human neuroblastoma cell lines induced by ATRA. We hypothesized a positive modulation of the antineoplastic effect of retinoids using LOX/COX inhibitors. Caffeic acid (3,4-dihydrocinnamic acid, CA), a phenolic plant compound exhibiting non-specific anti-oxidant effects both *in vivo* and *in vitro*, was chosen as a specific 5-lipoxygenase (5-LOX) inhibitor (25,26), and celecoxib (CX) was chosen as a cyclooxygenase-2 (COX-2) specific inhibitor (27-29).

Materials and methods

Cell lines. SH-SY5Y (ECACC cat. no. 94030304) and SK-N-BE(2) (ECACC cat. no. 95011815; *MYCN* amp.) neuroblastoma cell lines were used for this study. These cell lines were maintained in Dulbecco's modified Eagle's medium (DMEM)/Ham's F-12 medium mixture (1:1) supplemented with 20% fetal calf serum, 1% non-essential amino acids, 2 mM L-glutamine, 100 U/ml penicillin, and 100 mg/ml streptomycin (all from PAA Laboratories GmbH, Austria). Culture was performed under standard conditions at 37°C in a humidified atmosphere containing 5% CO₂. The cells were subcultured 1-2 times weekly.

Chemicals. ATRA (Sigma Chemical Co., St. Louis, MO, USA) was prepared as a stock solution at the concentration of 100 mM in dimethyl sulfoxide (DMSO; Sigma). CA (Sigma) and CX (LKT Laboratories, Inc., St. Paul, MN, USA) were dissolved in DMSO at the concentration of 130 and 100 mM, respectively. Reagents were stored at -20°C under light-free conditions.

Induction of differentiation. Stock solutions were diluted in fresh cell culture medium to obtain final concentrations of


1 and 10 μ M of ATRA, 13 and 52 μ M of CA and 10 and 50 μ M of CX. Concentrations of LOX/COX inhibitors were chosen on the basis of previously published data (25,30,31). The final concentration of DMSO in all experiments did not exceed 0.05% (v/v), and was found to influence neither cell proliferation nor induced differentiation. In all experiments, cells were seeded 24 h before the treatment, and untreated cells were used as a control.

Cell morphology. To examine cell morphology, 40,000 cells were seeded onto 8-cm² Petri dishes and allowed to attach for 24 h before treatment. Cell morphology was examined 2 and 7 days after ATRA and/or LOX/COX inhibitor treatment. Cell cultures were observed and recorded using an Olympus CKX41 inverted microscope in combination with an Olympus SP-350 camera.

Cell proliferation. The proliferation activity of cell populations, both treated and untreated, was measured by MTT and WST assay. For each experiment, cells were seeded in 96-well microtiter plates at a density of 10,000 cells per well and allowed to attach overnight. ATRA and/or LOX/COX inhibitors were added at the concentrations mentioned above. In order to perform the MTT assay, the culture medium was removed 48 h after treatment, and 220 μ l of DMEM/Ham's F-12 medium mixture (1:1) containing 3-[4,5-dimethylthiazol-2-yl]-2,5-diphenyltetrazolium bromide (MTT) (Sigma) at a final concentration of 455 μ g/ml in medium was added to each well. After a 4-h incubation under standard conditions, the culture medium with MTT was replaced by 200 μ l of DMSO per well to solubilize the MTT product. After a 10-min shaking of the microtiter plate, the absorbance was measured at 570 nm using a Sunrise Absorbance Reader (Tecan, Austria GmbH). To perform the WST assay, 10 μ l of Cell Proliferation Reagent WST-1 (Roche, Mannheim, Germany) was added to each well, and after a 4-h incubation under standard conditions and a 1-min shaking, the absorbance was measured at 450 nm with 620 nm reference wavelength using the Sunrise Absorbance Reader. Data from both MTT and WST assays were analyzed using two-way ANOVA followed by the Scheffé post-hoc test. $P < 0.001$ was considered significant.

Flow cytometry. To evaluate the cell cycle, 10,000 cells were seeded onto 25 cm² Petri dishes and allowed to attach overnight. The cells were harvested 3 and 10 days after short trypsinization and stained using Vindelov's solution [0.01 M Tris, 10 μ g/ml RNase, 50 μ g/ml PI (all from Sigma) and 0.1% Triton X-100 (ICN Biomedicals, Irvine, CA, USA) and 1 mM NaCl] for 30 min at room temperature (RT). The FACS Canto™ II flow cytometer with I BD FASC DIVA Software (Beckton Dickinson, CA, USA) was employed to analyze the cell cycle at 3 and 10 days after treatment. Twenty thousand events per sample were evaluated using WinMDI 2.8 software (Joseph Trotter, Scripps Research Institute, La Jolla, CA, USA).

Antibodies. To perform the Western blot analysis, the following primary antibodies were used: rabbit monoclonal anti-glial fibrillary acidic protein (GFAP) antibody (cat. no. G4546, Sigma), mouse monoclonal anti-neurofilament 200

 SPANDIDOS PUBLICATIONS antibody (cat. no. N0142, Sigma), mouse mono-
anti-synaptophysin antibody (cat. no. S5768, Sigma),
mouse monoclonal anti-neuronal nuclei (NeuN) antibody
(cat. no. MAB377, Millipore, Germany) and mouse mono-
clonal anti-cellular retinoic acid-binding protein 1 (CRABP 1)
antibody (cat. no. Ab2816, Abcam, Cambridge, UK).
Secondary antibodies for immunoblotting included anti-
mouse IgG antibody conjugated with peroxidase (cat. no.
A9917, Sigma) and anti-rabbit IgG antibody conjugated with
peroxidase (cat. no. A2074, Sigma). As a control, mouse
monoclonal anti- α -tubulin (cat. no. T5168, Sigma) was used.

Immunoblotting. Whole-cell extracts were loaded onto polyacrylamide gels, electrophoresed, and blotted onto polyvinylidene difluoride membranes (PVDF, Bio-Rad Laboratories GmbH, Germany). The membranes were blocked with 5% nonfat milk in phosphate-buffered saline with 0.1% Tween 20 (PBS-T), and incubated overnight at 4°C with selected antibodies diluted 1:1,000 in blocking solution. After rinsing with PBS-T (at least three 10-min washes), membranes were incubated with secondary antibody at 1:10,000 dilution for 45 min at RT. Each step was followed by at least three 10-min washes in PBS-T. ECL-Plus detection was performed according to the manufacturer's instructions (Amersham, GE Healthcare, UK).

Results

Our experiments were focused on a detailed study of the possible effect of combined treatment with ATRA and LOX/COX inhibitors (CA or CX) on neuroblastoma cells. All experiments were performed using the compounds mentioned above alone and in combinations. The course of cell differentiation was evaluated using cell morphology assessment and detection of intracellular markers by immunoblotting. The bioavailability of ATRA in the treated cell lines was studied using detection of CRABP-I protein by immunoblotting. The proliferation activity of treated cells was measured using MTT and WST assays. Changes in cell cycle and the proportion of apoptotic cells were evaluated by the use of flow cytometry.

Cell differentiation. During long-term culture, the morphology of both neuroblastoma cell lines markedly changed after induction of differentiation by ATRA, either alone or in combination with LOX/COX inhibitors in defined concentrations.

The changes in cell morphology were observed within 7 days after treatment, when the control untreated cells reached confluence. After treatment with ATRA, the first morphological changes were detectable at day 2, and they were most pronounced at day 7 (Fig. 1). ATRA applied alone (at both 1 and 10 μ M concentrations) or in combination with CA (at both 13 and 52 μ M concentrations), reduced cell proliferation and induced marked changes in cell morphology comprising the appearance of long cytoplasmic protrusions and a general neuron-like phenotype. Concerning the combination of ATRA with CA, more obvious changes in cell morphology were observed in SK-N-BE(2) cells; therefore, this cell line seems to be more sensitive to ATRA treatment

in general (Fig. 1C-J). On the contrary, CA applied alone at 13 or 52 μ M concentrations did not affect the differentiation of both cell lines. In the presence of CA, cells proliferated rapidly to confluence (day 7 after treatment), and their morphology did not change.

Treatment of both neuroblastoma cell lines with 10 or 50 μ M CX in combination with 1 or 10 μ M ATRA also led to differentiation into neuron-like phenotypes, particularly in combination with 10 μ M ATRA (Fig. 1S-V). However, CX applied either alone or in combination with ATRA showed a cytotoxic effect that led to an observable decrease in total cell number in culture. This phenomenon was explicitly detectable in the SH-SY5Y cell line (Fig. 1Q, R, U and V).

The course of cell differentiation in both cell lines treated with ATRA and/or LOX/COX inhibitors was also investigated in detail using immunodetection of specific differentiation markers (Fig. 2).

The induction of neuronal differentiation was confirmed by detection of neurofilament 200 (NF-200) expression. To investigate the potential ability of ATRA to induce glial differentiation, GFAP expression levels were also examined. To further verify the induction of neuronal differentiation in neuroblastoma cell lines, expression of NeuN as a marker of terminal differentiation of neuronal cells and synaptophysin as a neuroendocrine tumor marker was evaluated.

The expression levels of NF-200 and NeuN increased in the ATRA-treated cells of both cell lines within 7 days after treatment, confirming the potential of ATRA to induce neuronal differentiation in both neuroblastoma cell lines (Fig. 2). On the other hand, conflicting results were obtained for detection of synaptophysin: while the expression levels of synaptophysin obviously increased in both ATRA-treated and untreated SK-N-BE(2) cells in a concentration-dependent manner, the situation in the SH-SY5Y cell line was exactly the opposite (Fig. 2). In addition, the differentiation into neuronal phenotype was confirmed indirectly by complete absence of GFAP (data not shown). Furthermore, the treatment of both cell lines with LOX/COX inhibitors alone had no influence on the differentiation process since the expression of all markers was the same as in the control cells (data not shown).

The application of ATRA in combination with CA led to partly inverted results in the neuroblastoma cell lines. The enhancement of ATRA-induced neuronal differentiation by combined application with CA was confirmed in both cell lines when NF-200 was used as a marker; moreover, this enhancement by CA showed concentration dependence (Fig. 2). However, the expression of NeuN upon treatment with ATRA and CA combinations was slightly increased in SK-N-BE(2) cells and markedly decreased in SH-SY5Y cells at day 7 after treatment, especially when 1 μ M ATRA was used. Furthermore, a completely inverted pattern of synaptophysin expression was found in these cell lines after the same combined treatments with ATRA and CA (Fig. 2).

Detection of the same differentiation markers in both neuroblastoma cell lines after application of CX in combination with ATRA mainly corresponded to results obtained from experiments with CA/ATRA combinations. Increased levels of NF-200 after ATRA and CX combined treatment were observed in the early phases of the induced

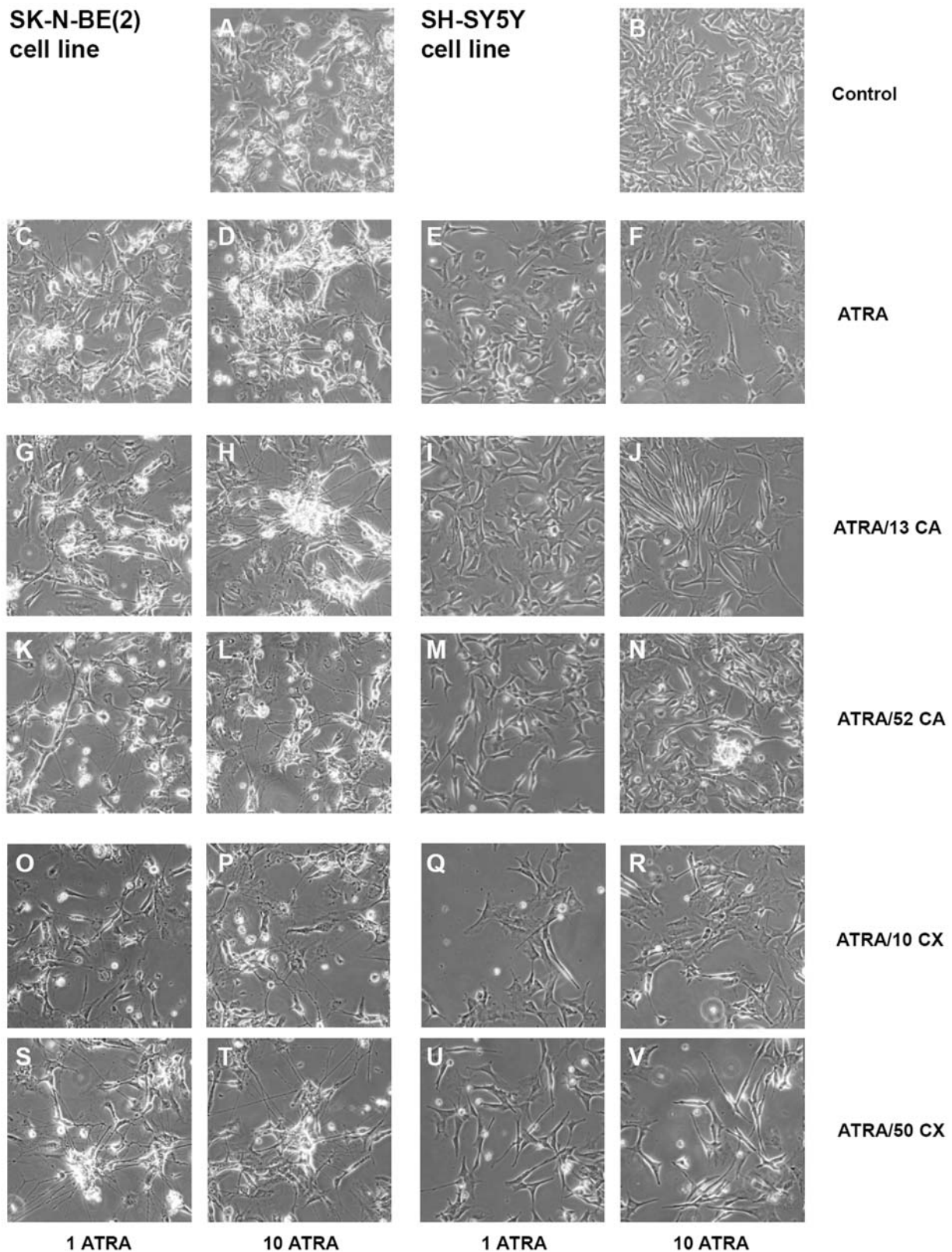


Figure 1. Morphology of SK-N-BE(2) and SH-SY5Y neuroblastoma cells, both untreated (A and B) and treated with ATRA alone (C-F) in combinations with CA (G-N) or CX (O-V). ATRA was applied in concentrations of 1 or 10 μM (1 ATRA, 10 ATRA); CA in concentrations of 13 and 52 μM (13 CA, 52 CA), and CX in concentrations of 10 and 50 μM (10 CX, 50 CX). Micrographs were taken at day 7 after treatment. Original magnification, $\times 100$.

differentiation. However, results obtained from the late phases of the cell differentiation were more inconsistent, probably due to the cytotoxic effect of CX as documented in the cell morphology experiments (Figs. 1O-V and 2). The

same inverse expression pattern of NeuN in both cell lines, i.e., the increase in SK-N-BE(2) cells and decrease in SH-SY5Y cells, was detected after combined treatment with ATRA and CX, similarly to the combinations with CA

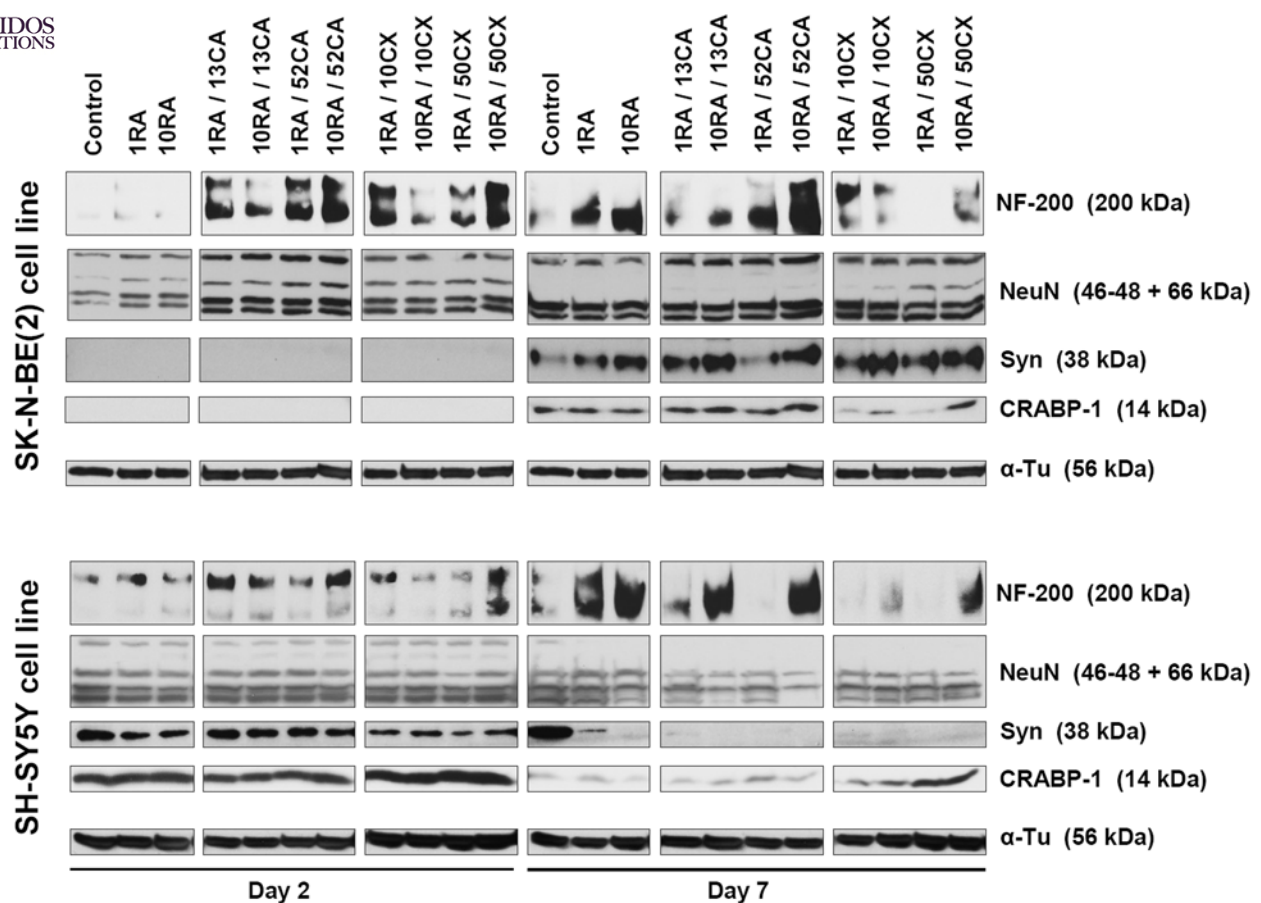


Figure 2. Western blot analysis of differentiation markers and CRABP-1 expression in SK-N-BE(2) and SH-SY5Y cells, both untreated and treated with ATRA alone or in combinations with CA or CX. ATRA was applied in concentrations of 1 or 10 μM (1RA, 10RA); CA in concentrations of 13 and 52 μM (13CA, 52CA), and CX in concentrations of 10 and 50 μM (10CX, 50CX). The immunoblotting was performed at days 2 and 7 after treatment.

described above. The same inverse pattern of synaptophysin expression was found in both cell lines after ATRA/CX combined treatment and after ATRA/CA combined treatment (Fig. 2). Moreover, changes in the spectrum of detectable NeuN bands were apparent: while the 48-kDa band faded away in a time-dependent manner in SK-N-BE(2) cells (except for those treated with ATRA/CX), the 66-kDa band was absent in SH-SY5Y cells after 7 days of treatment.

To evaluate the intracellular activity of ATRA, expression levels of CRABP-1 were investigated simultaneously. In general, the expression patterns of CRABP-1 in both cell lines were in accordance with the inverse expression pattern observed for synaptophysin: CRABP-1 levels in SK-N-BE(2) cells increased during the cell differentiation and exhibited a slight concentration dependence after combined treatments. In the SH-SY5Y cell line, the situation was completely inverse, i.e., CRABP-1 expression markedly decreased in the late phase of cell differentiation, and only combined treatment with 50 μM CX resulted in maintained expression (Fig. 2).

Cell proliferation. The influence of all treatments used on the proliferation activity of both selected cell lines was examined by MTT assay for experiments with CA and combinations with ATRA, or by WST assay for CX and its combinations. The data are expressed as relative mean absorbance (ratio of treatment/control).

The application of CA alone, at either 13 or 52 μM concentrations, generally did not affect the proliferation activity of both of these cell lines (Fig. 3A and B). Treatment with ATRA alone significantly stimulated proliferation of SK-N-BE(2) cells; however, neither stimulation nor inhibition was observed for any other treatment in this cell line (Fig. 3A). In the SH-SY5Y cell line, a slight anti-proliferative effect was detectable after combined treatment with 10 μM ATRA with CA at both concentrations (Fig. 3B).

A strong cytotoxic effect of CX alone was confirmed by WST assay in both of these cell lines (Fig. 3C and D); these data are in accordance with the results obtained with cell morphology as described above. On the other hand, the application of ATRA alone at both concentrations did not significantly change the proliferation activity of SH-SY5Y cell line (Fig. 3D). However, the combined treatment led to different results: while the proliferation activity of SH-SY5Y cells was diminished by CX application independently of its combination with various concentrations of ATRA (Fig. 3D), the combined treatment with CX and ATRA reduced the inhibitory effect of CX in a concentration-dependent manner in SK-N-BE(2) cells (Fig. 3C).

Cell cycle and apoptosis. Finally, flow cytometry was employed to investigate in detail the influence of combined treatment with ATRA and the LOX inhibitor CA on cell

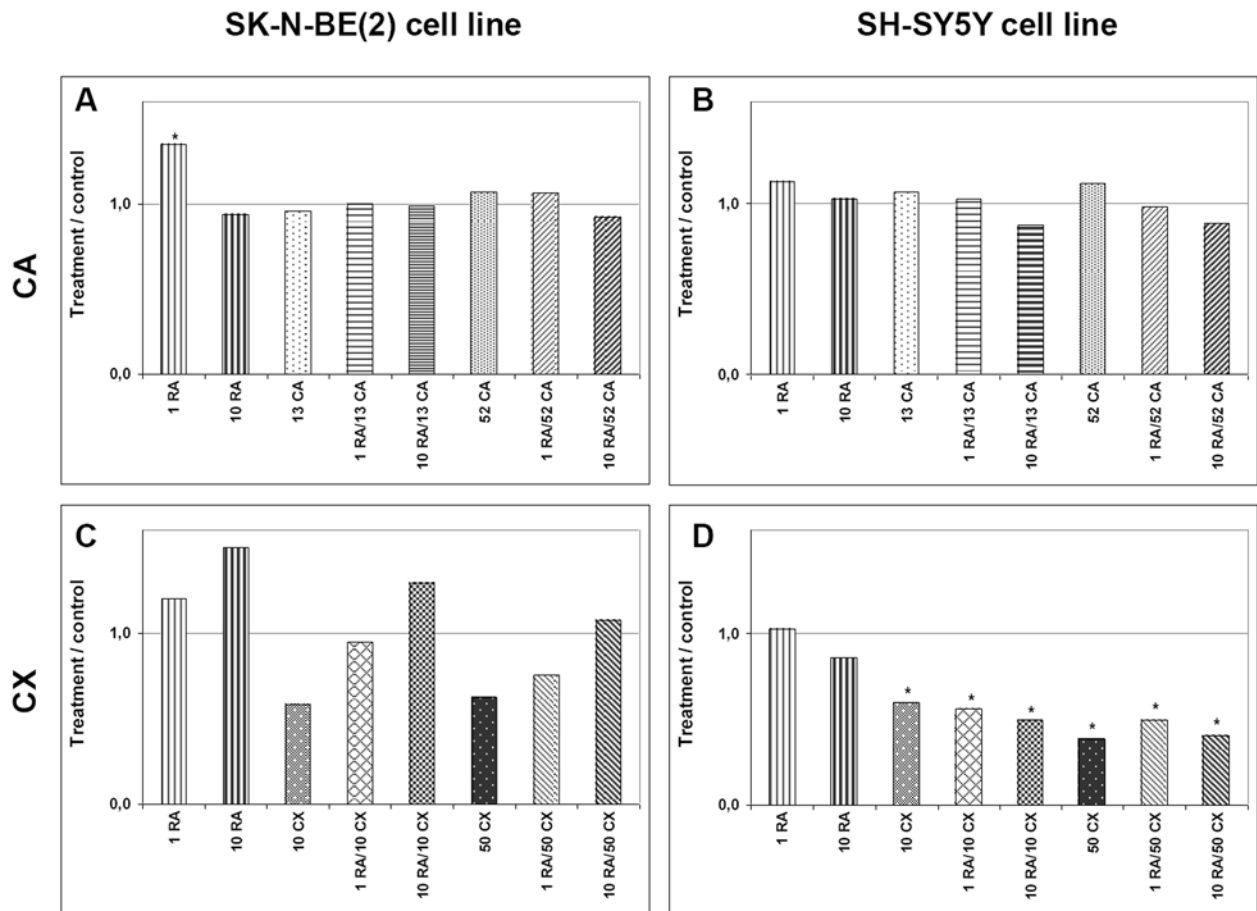


Figure 3. Proliferation activity of SK-N-BE(2) and SH-SY5Y cells treated with ATRA and/or LOX/COX inhibitors measured by MTT assay (A and B) and WST-assay (C and D), respectively. X-axis, treatment; Y-axis, relative cell number (ratio of treatment/control). Data were analyzed using two-way ANOVA followed by the Scheffé post-hoc test. *Significant difference ($P < 0.001$) from the control group.

cycle and apoptosis induction in selected neuroblastoma cell lines. Application of CA alone changed neither the cell cycle course nor apoptosis frequency in SK-N-BE(2) cells (Table I); however, this compound exhibited a moderate pro-apoptotic effect accompanied by changes in the cell cycle in SH-SY5Y cells (Table II). In accordance with our other results described above, flow cytometric analysis of the DNA content confirmed higher sensitivity of the SK-N-BE(2) cell line to the combined treatment upon comparison to SH-SY5Y cells (Tables I and II, Fig. 4).

Combined application of CA with 1 μM ATRA enhanced the differentiation of SK-N-BE(2) cells in a concentration-dependent manner: an increase in the number of cells in the G_1 -phase accompanied by a decrease in the number of cells in the G_2/M phase was detected (Table I, Fig. 4A and C). In contrast, application of CA in combination with 10 μM ATRA led to the shift of peak positions indicative of an increase in the number of cells in G_2/M phase; in particular, an increase in the sub-diploid peak was observed which corresponded to a higher frequency of apoptosis (Table I, Fig. 4E and G).

Slight changes in the cell cycle course corresponding to induced differentiation, i.e. an increase in the number of cells in the G_1 -phase and a decrease in the number of cells in the G_2/M phase, were observed in SH-SY5Y cells when 52 μM CA was used for combined treatment with both concentrations

of ATRA (Table I, Fig. 4D and H). Combined treatment with 13 μM CA did not induce any similar effect in the same cell line (Table I, Fig. 4B and F). Furthermore, the use of 10 μM ATRA alone led to the shift of the G_2/M peak in the SH-SY5Y cells.

Discussion

Neuroblastoma is the most frequent extracranial solid tumor in children. It is considered to be an embryonic tumor arising from immature cells of the neural crest, probably due to genetic and molecular alterations that result in arrested cell differentiation and uncontrolled proliferation. In general, neuroblastic tumors are characterized by extreme clinical and pathological heterogeneity (32). The induced differentiation of tumor cells is regarded to be therapeutically advantageous, since more extensively differentiated neuroblastic tumors are usually associated with lower stage and better clinical outcome. Therefore, differentiating agents such as retinoic acid, which is known to induce differentiation in several tumor types including neuroblastoma, have become a part of the therapeutic arsenal (10-12,20,32).

The biological effects of RA and retinoids are generally mediated by inducible nuclear retinoid receptors (RARs and RXRs), which regulate transcription of genes with RA-

SPANDIDOS PUBLICATIONS Flow cytometric analysis of the cell cycle in SK-N-BE(2) cells.^a

Treatment	% <G ₁	% G ₁	% S	% G ₂ /M
Control	10.9	74.6	11.6	2.9
13 CA	13.2	70.5	10.8	5.5
52 CA	10.0	73.0	11.8	6.2
1 ATRA	35.6	29.0	21.9	13.5
1 ATRA/13 CA	38.0	25.2	20.9	15.9
1 ATRA/52 CA	33.2	37.7	14.9	14.2
10 ATRA	30.4	49.4	10.0	10.2
10 ATRA/13 CA	37.3	25.7	15.2	21.8
10 ATRA/52 CA	34.7	33.2	13.9	18.2

^aData from the representative experiment. Cells were analyzed at day 10 after treatment. ATRA was applied in concentrations of 1 or 10 μ M (1 ATRA, 10 ATRA); CA in concentrations of 13 and 52 μ M (13 CA, 52 CA).

Table II. Flow cytometric analysis of the cell cycle in SH-SY5Y cells.^a

Treatment	% <G ₁	% G ₁	% S	% G ₂ /M
Control	40.1	25.4	16.5	18.0
13 CA	35.4	26.0	19.4	19.2
52 CA	37.0	24.9	19.3	18.8
1 ATRA	47.6	22.2	18.1	12.1
1 ATRA/13 CA	45.5	22.5	17.2	14.8
1 ATRA/52 CA	44.2	20.3	18.9	16.6
10 ATRA	55.4	20.7	11.9	12.0
10 ATRA/13 CA	54.5	18.5	15.7	11.3
10 ATRA/52 CA	43.7	19.8	18.4	16.1

^aData from the representative experiment. Cells were analyzed at day 10 after treatment. ATRA was applied in concentrations of 1 or 10 μ M (1 ATRA, 10 ATRA); CA in concentrations of 13 and 52 μ M (13 CA, 52 CA).

responsive elements (32,33). The cellular retinol-binding proteins (CRBP) and cytoplasmic retinoic acid-binding proteins (CRABP-1 and CRABP-2) enhance the binding of RA to their respective receptors (34). Despite the beneficial clinical effects of treatment with RA, as mentioned above, evidence of intrinsic or acquired resistance, as well as of potential toxicity, considerably limit the use of retinoids in clinical practice. Strategies to overcome these problems consist of efforts to reduce toxicity, retain bioavailability, and develop effective combined curative regimens (35-37).

The aim of our work was to examine the possible potentiation of retinoic acid-induced differentiation of SH-SY5Y and SK-N-BE(2) cells by inhibition of intracellular degradation of ATRA. Although Reynolds and co-workers

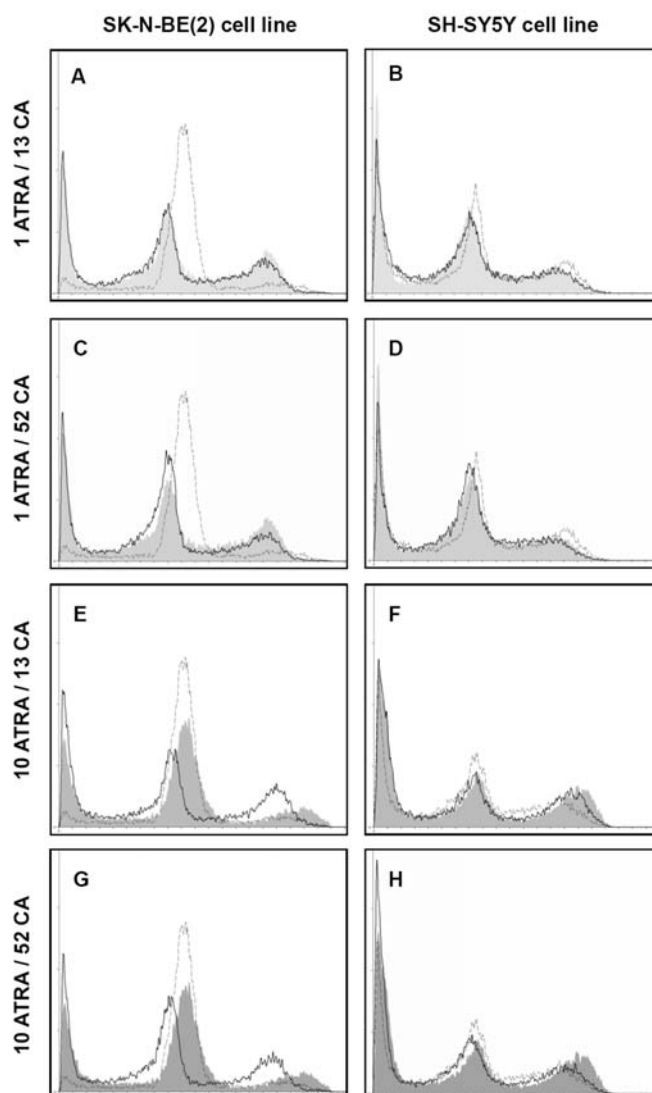


Figure 4. Representative flow cytometric analysis of the DNA content in SK-N-BE(2) and SH-SY5Y neuroblastoma cell lines at day 10 after treatment. ATRA was applied in concentrations of 1 or 10 μ M (1 ATRA, 10 ATRA); CA in concentrations of 13 and 52 μ M (13 CA, 52 CA). Histograms show results for untreated cells (open peak) overlaid with cells treated with ATRA alone (gray) and with combinations of ATRA and CA (continuous line). X-axis, logarithm of fluorescence (DNA content); Y-axis, number of events.

(20) preferred 13-*cis*-RA for neuroblastoma treatment, we decided to use ATRA to induce cell differentiation based on the conclusions of Armstrong and colleagues (21) that all isomers of retinoic acid undergo intracellular conversion to ATRA. CA, as an inhibitor of 5-LOX, and CX, as an inhibitor of COX-2, were chosen to study the possibility of the modulation of ATRA-induced cell differentiation in neuroblastoma cells. We investigated cell differentiation, cell proliferation, cell cycle and apoptosis in two neuroblastoma cell lines treated either with ATRA alone or with ATRA combined with LOX/COX inhibitors.

After treatment with ATRA, marked changes in cell morphology and an increased number of cells exhibiting a neuron-like phenotype were detectable in both cell lines within 7 days; this effect was more obvious in SK-N-BE(2) cells. This observation is in accordance with results obtained with

other methods that all confirmed higher sensitivity of SK-N-BE(2) cells to ATRA-induced differentiation, as well as enhancement by combined treatment with CA. Analysis of cell morphology also indicated a potentiation of ATRA-induced differentiation by application of selected inhibitors in these cell lines. While treatment with inhibitors alone did not lead to the neuron-like phenotype, marked changes in cell morphology were detectable after application of ATRA alone or in combination with the inhibitors. Moreover, a strong cytotoxic effect of CX was observed in both of these cell lines.

To investigate the course of induced differentiation in neuroblastoma cell lines in detail, expression levels of specific differentiation markers were monitored. The course of neuronal differentiation in both ATRA-treated neuroblastoma cell lines was confirmed by detection of NF-200, NeuN, and synaptophysin; expression of GFAP was not detected at all. Monitoring of the expression levels of the listed markers revealed that ATRA has the potential to induce neuronal differentiation in the selected neuroblastoma cell lines, as previously described in the literature (38,39). However, some ambiguous results were obtained for detection of synaptophysin expression levels: while SK-N-BE(2) cells exhibited a shift to neuronal phenotype that was time- and concentration-dependent, the situation in SH-SY5Y cells was completely inverse.

A similar discrepancy in the expression pattern of differentiation markers between these two cell lines was also observed after combined application of ATRA with inhibitors. In general, although LOX/COX inhibitors applied alone had no impact on the course of differentiation, their capability to modulate and further enhance ATRA-induced differentiation was confirmed in the SK-N-BE(2) and SH-SY5Y cell lines when NF-200 and NeuN were employed. This partial inconsistency in results for SH-SY5Y cells treated with ATRA in combination with CX was probably caused by the cytotoxicity of this compound, which also prevailed over cell differentiation in morphology studies.

The partially inverted pattern of results for SK-N-BE(2) and SH-SY5Y cells was probably caused by the different tumor cell type that these cell lines represent: while the SH-SY5Y cell line belongs to the N-type of neuroblastoma cell lines, SK-N-BE(2) cell line exhibits the characteristics of intermediate I-type (40,41). In general, different biological features (cell shape, nuclear/cytoplasmic ratio, adherence to the substrate, differentiation potential, tumorigenicity) of N-, I- and S-types of neuroblastoma cell lines are independent on neither *MYCN* amplification status nor amount of N-myc protein. The I-type was proved to exhibit a stem cell phenotype with higher differentiation potential and sensitivity to RA (41). The stem cell phenotype of SK-N-BE(2) cells was also recently confirmed by analysis of specific stem cell markers (42). Our results for SK-N-BE(2) and SH-SY5Y cell lines are in accord with these findings. Moreover, the described differences in N- and I-types of neuroblastoma cell lines can explain the distinctions between cell lines observed in our study.

The results from the flow cytometric analysis also demonstrated higher sensitivity of SK-N-BE(2) cells to the combined treatment with ATRA and CA. The enhancement

of induced differentiation was confirmed after combined treatment with 52 μ M CA and 1 μ M ATRA in SK-N-BE(2) cells, whereas a moderate pro-apoptotic effect was obvious after combined treatment of CA and 10 μ M ATRA in these cells, as well as in SH-SY5Y cells. The increase in the sub-diploid cell fraction corresponding to a higher frequency of apoptosis was relatively surprising, since the published data suggest that ATRA does not induce apoptosis in neuroblastoma cells (35,37,43).

The expression levels of CRABP-1 were investigated simultaneously to evaluate ATRA intracellular activity and availability. CRABP-1 is one of the contributors to the regulation of RA signaling pathways, sharing 75% amino acid homology and having the same molecular weight as CRABP-2 (44,45). The presence or absence of different combinations of RA-binding proteins is a critical determinant of the cellular response to RA (46). The functions of CRABPs have not yet been decoded, and CRABP-1 may function differently from CRABP-2. The one feature they clearly have in common is their capability to serve as a shuttle to move RA through the aqueous cytosol to the nucleus, where RA can interact with RARs and initiate transcription of various genes. Kinetic studies of the movement of RA to RARs showed that while CRABP-1 is a passive vehicle, binding and releasing its ligand depending on concentration gradients, CRABP-2 can deliver RA to RARs in a direct collisional process (47). The detected pattern of CRABP-1 expression correlates with the observed trends in synaptophysin expression: while slight concentration- and time-dependent characteristics of CRABP-1 expression were apparent in SK-N-BE(2) cells, the results in SH-SY5Y cells were inverted. The only exception was the combined treatment with 50 μ M CX after 7 days, which also maintained CRABP-1 expression in SH-SY5Y cells.

It has been reported that CRABP-1 and RA are inversely regulated (48). Furthermore, CRABP-1 binds RA and prevents it from entering the nucleus; in cells with low CRABP-1 expression, RA enters the nucleus (49-51). Uhrig and colleagues (52) demonstrated that up-regulation of CRABP-1 reduced the differentiation potential of SH-SY5Y cells. However, all these findings were concerned with the physiological plasma concentration of RA in humans, which is approximately 10 nM (53). Excess exogenous RA may oversaturate the binding capacities of CRABP-1, allowing the remaining RA to bind to RARs (54). For this reason, the inverse expression patterns of CRABP-1 observable in SK-N-BE(2) and SH-SY5Y cells may have been partly caused by the differences in cell type as mentioned above, and partly by the differential expression pattern of RA receptors across cell types.

Results of cell proliferation assays also confirmed the differences in biological behavior for these cell lines according to their sensitivity to the applied treatment. Even ATRA treatment alone showed different responses for these cell lines: while the proliferation activity of SK-N-BE(2) cells was significantly stimulated by application of ATRA alone, the same treatment in SH-SY5Y cells did not significantly increase the proliferation activity. It has been reported that treatment with ATRA results in a concentration-dependent decrease in cell proliferation (39) but other authors have



enhanced cell viability after application of RA the contrary, the response of both investigated cell lines to treatment with LOX/COX inhibitors applied alone was similar: CA applied alone did not affect proliferation, whereas application of CX alone revealed a strong cytotoxic effect. In contrast to results obtained for the course of cell differentiation, the SH-SY5Y cell line seemed to be more sensitive, exhibiting a slight anti-proliferative effect when CA was applied in combination with 10 μ M ATRA; a similar effect was not apparent in SK-N-BE(2) cells. However, the proliferation activity reflects the course of cell differentiation only indirectly, and this effect is probably caused by the induction of apoptosis, as apparent from the result obtained with flow cytometry in this cell line. Also, the combined treatment with CX led to different results for both of these cell lines: while the combined CX/ATRA treatment in SH-SY5Y cells decreased the proliferation activity independently of the concentrations used, in SK-N-BE(2) cells the combination of CX and ATRA reduced the inhibitory effect of CX alone in a concentration-dependent manner. Although we originally supposed that the reduction of proliferation activity in neuroblastoma cell lines is associated with the process of induced differentiation as already published by other authors (34,38), the obtained results also suggest the possibility of apoptosis induction by combined treatment with ATRA and inhibitors, or a direct cytotoxic effect of CX that may be partly diminished by combined application with ATRA.

The design of our study was based on the hypothesis that inhibition of 5-LOX and/or COX-2 pathways in arachidonic acid metabolism may enhance the course of cell differentiation after treatment with ATRA in neuroblastoma cells. Our results for the SK-N-BE(2) cell line support this hypothesis, especially when CA was used as the inhibitor of 5-LOX. A similar effect of CA and its phenethyl ester on ATRA-induced cell differentiation was already shown for the HL-60 leukemia cell line (25,56). This differentiation enhancement might be explained by imbalances in arachidonic acid metabolism after application of LOX/COX inhibitors. These imbalances probably increase the intracellular amount of free arachidonic acid or arachidonic acid metabolites that can potentiate the differentiation effect of RA; similar results were documented after application of exogenous arachidonic acid (24). However, this modulation of cell differentiation was less apparent when CX was used as the inhibitor of COX-2, particularly due to its cytotoxic effect, which may prevail over the possible influence of CX on ATRA-induced cell differentiation.

In conclusion, our results suggest a distinct response of SK-N-BE(2) and SH-SY5Y cells to the combined treatment. The SK-N-BE(2) cell line, which is considered to be of stem cell phenotype, was noticeably more sensitive to the enhancement of ATRA-induced cell differentiation. These results suggest that cancer stem cells, or rather tumor-initiating cells, might be the most suitable target for enhanced cell differentiation. Nevertheless, more detailed investigation of the possible mechanisms of combined induced differentiation with regard to differences in neuroblastoma cell type are needed. Therefore, our forthcoming study is focused on gene expression profiling of both of these neuroblastoma cell lines after combined treatment with ATRA and LOX/COX inhibitors.

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