Abstract. Rat C6 glioma is a chemo-resistant experimental brain tumor that is difficult to treat with various drug combinations. Previous studies suggested that imatinib mesylate (Gleevec) is effective in pre-clinical trials for glioblastoma. Also, chlorimipramine (Anafranil) is an anti-depressant drug in use in the clinic and shown to have anti-neoplastic activity. We hypothesized that treatment of resistant C6 glioma with combination of imatinib and chlorimipramine may potentiate cytotoxicity and reverse resistance. C6 glioma was examined both as monolayer and as spheroid cultures. Several experimental designs were examined all of which showed synergistic activity albeit at different time kinetics. Combination treatment resulted in inhibition of cell growth and enhanced cell death as determined by dye exclusion. Further, the combination treatment resulted in significant induction of apoptosis as determined by Annexin V-FITC and PI. Also, there was inhibition of DNA synthesis and cAMP. Altogether, these findings supported the anti-proliferative and cytotoxic effects of the combination treatment. Morphological studies were also performed using transmission and scanning electron microscopy. Significant synergistic apoptosis was detected by the combination treatment in both the monolayers and spheroid cultures. There was also a synergistic effect in autophagy by the combination. Several altered morphological features were noted by both the individual compound and enhanced by the combination treatment. The present findings support our hypothesis and demonstrate the potentiation of cytotoxicity by the combination of imatinib and chlorimipramine in C6 glioma. Further, the findings suggest the potential clinical application of the combination in the treatment of drug-resistant glioma.

Introduction

Receptor and non-receptor tyrosine kinases (TKs) have emerged as clinically useful drug target molecules for treating cancer (1). Imatinib mesylate (IM; STI-571, Gleevec®) is a small molecule ATP analogue which is a potent and selective inhibitor of several protein TKs of the receptors for platelet-derived growth factor (PDGF), bcr-abl, stem cell factor (SCF), and c-kit by interacting with the ATP-binding site (2,3). It has been reported that mitochondria and telomerase are also targets of imatinib mesylate (4,5). IM was primarily designed to treat chronic myeloid leukemia (CML), but it is also currently the drug of choice for the treatment of metastatic gastrointestinal stromal tumors (GISTs) (6,7). The success of this compound in the treatment of CML and GISTs has led to the broader examination of its application in the treatment of other tumors, such as glioblastoma (8), small lung cancer (9), anaplastic thyroid cancer (10) and prostate cancer (11).

C6 rat glioma is a chemo-resistant experimental brain tumor which is difficult to treat with a combination of drugs (12). This cell line has properties similar to human glioblastoma and has been widely used as a model system for investigation of human glioblastoma (13). Imatinib has recently been found to be efficacious in preclinical trials for glioblastoma, but the use of IM has been tempered by the problems of disease persistence and resistance. It has been reported that IM inhibits the growth of glioma cells in preclinical models, but its poor brain distribution by P-glycoprotein (P-gp) limits its efficacy in patients (14,15).
Chlorimipramine (CIMP; clomipramine or Anafranil®) is the tricyclic anti-depressant that has been in clinical use for over 30 years (16). Previous studies which were made on various types of cancer cells, including drug-resistant leukemias, renal cancer cells, solid murine tumors, and a wide range of cell cultures derived from brain tumors, (e.g., astrocytomas, mixed oligo-astrocytomas, glioblastoma multiforme, and meningiomas), revealed the novel effect of CIMP as an anti-neoplastic agent (17-20). From studies carried out on non-tumorigenic tissues and tumorigenic cell lines, it has been shown that CIMP inhibits complex-III of the respiratory chain, resulting in increased ROS, cytochrome C release and caspase-activated apoptosis (21,22). CIMP also causes irreversible DNA damage in both normal and neoplastic cells, but mitochondria damage has only been observed in neoplastic cells (19). CIMP has been demonstrated to reverse multidrug resistance (MDR) to conventional antineoplastic agents, such as doxorubicin, in several tumor cell lines in vitro (17,19,23,24).

One of the most recent novel developments in chemotherapy has been to move away from cytotoxic agents that inhibit non-specific targets in cellular pathways that contribute to carcinogenesis and resistance (25,26). The present study investigated the therapeutic potential of the selective mitochondrial inhibitor, CIMP, used in combination with IM for the treatment of malignant C6 glioma cells used as monolayers and spheroid cultures. This combination of IM and CIMP has not been previously investigated in vitro or in vivo.

Materials and methods

Drugs and solutions. The following drugs were used: IM was kindly provided by Dr E. Buchdunger (Novartis AG, Basel, Switzerland) and Dr Fulya Dincturk (Novartis AG, Istanbul, Turkey). CIMP was obtained from Boehringer (Mannheim, GmbH). IM was prepared as 100 μM stock solution in phosphate buffered saline (pH 7.2) and filtered through a 0.22 μ injector filter (TPP, Trasadingen, Switzerland) before use. CIMP was dissolved in 70% ethanol to give a stock solution of 100 μM.

Monolayer culture. The rat C6 glioma cell line was supplied by ATCC (Rockville, MD, USA) and was grown in monolayers in Dulbecco’s Modified Eagle’s Medium-F12 (DMEM-F12; Biological Industries, Haemek, Israel) supplemented with 10% heat-inactivated foetal calf serum (FCS; Sigma Chemical Co., St. Louis, MO, USA), 100 units/ml penicillin and 100 μg/ml streptomycin (both from Sigma Chemical Co.). Cells in semi-confluent flasks were harvested by using 0.05% trypsin (Sigma Chemical Co.), centrifuged after adding DMEM-F12 for trypsin inactivation, and resuspended in DMEM-F12. After trypsin blue exclusion, the glioma cells were plated in 6-well culture plates containing 5 ml DMEM-F12 medium at a concentration of 5x10^5 cells/well with 100% viability.

Constitution of a 3-dimensional spheroid model. An in vitro multicellular C6 glioma spheroid model was established using the liquid overlay technique. Briefly, once the monolayer cultures became semi-confluent, the cells were trypsinized and single cells with 100% viability were cultured over 3% Difco Noble agar-coated (Becton-Dickinson, Sparks, MD) 6-well culture plates containing 5 ml DMEM-F12 medium at a concentration of 1x10^5 cells/well (27).

Experimental design. The investigated drugs at a volume of 100 μl were applied to monolayers and spheroid cultures of malignant C6 glioma cells at the following final concentrations: 1, 10, 100 and 200 μM, except for the control cells, to which nutrient medium was applied. The cultures were incubated for 24, 48, 72 and 96 h. The IC50 values of IM and CIMP were determined as 10 μM. Four sets of experiments were performed under the following standard monolayer and spheroid culture conditions: i) untreated control cells; ii) cells treated with IM; iii) cells treated with CIMP; and iv) cells treated with a combination of IM and CIMP at a ratio of 1:1. The effects of drugs, both on monolayers and spheroid cultures, were determined by evaluation of cytotoxicants, including cell proliferation (total cell number), cell viability by trypan blue exclusion assay, the bromodeoxyuridine labeling index (BrDU-LI), spheroid volumes, morphology by transmission (TEM) and scanning (SEM) electron microscopy, Annexin V-FITC/PI staining, and cAMP levels by radioimmunoassay.

Cell cycle kinetics

Analysis of DNA synthesis (S-phase) by BrdU incorporation for monolayer cultures. C6 rat glioma cells were seeded on the cover slips and treated with drugs, as described in the survival studies throughout 96 h at 24-h intervals. Endogenous peroxidase activity was quenched with 0.5% H2O2 in methanol. Double-stranded DNA was denatured with 4 N HCl. Following the washings, a non-specific blocking reagent (Ultra-V-Block; Lab Vision Co., Westhinghouse, CA, USA) was used to prevent non-specific binding. Monoclonal mouse anti-BrdU antibody (1:50; Lab Vision Co.) was the primary antibody and the secondary antibody was biotinylated goat anti-mouse antibody (Lab Vision Co.). After washing, peroxidase-conjugated streptavidin (Lab Vision Co.) was applied and aminoethyl carbazole was used as a chromogen. The cover slips were then counterstained with Mayer's hematoxylin to augment the nuclear staining. BrDU-labelled cells in the S-phase had red-stained nuclei. For negative controls, the cover slips of the same groups were processed with exactly the same steps, excluding the primary antibody. The BrDU-LI (the number of positively stained cells divided by the counted total number of cells) was calculated by evaluating at least 3000 cells in multiple high power fields.

DNA synthesis (S-phase) analysis by BrdU incorporation for spheroid cultures. Spheroids were gently removed from the surface of the solidified agarose. The sections were treated with 0.1% trypsin for antigen retrieval and incubated in a solution of 0.3% H2O2. The sections were treated with 4 N HCl and incubated in 0.1 M Borax. Following the washings, a non-specific blocking reagent (Ultra-V-Block; Lab Vision Co.) was used to prevent non-specific binding. Monoclonal mouse anti-BrdU antibody (1:50; Lab Vision Co.) was the
primary antibody and the secondary antibody was a biotinylated goat anti-mouse antibody (Lab Vision Co.). After washing, peroxidase-conjugated streptavidin (Lab Vision Co.) was applied and aminoethyl carbazole was used as a chromogen. The sections were counterstained in Mayer’s hematoxylin. BrDU-labelled cells in S-phase had red-stained nuclei. Approximately 30 spheroids in each slide were determined for the calculation of the BrDU-LI. For negative controls, adjacent sections were processed with exactly the same steps, excluding the primary antibody.

Annexin V-FITC/PI staining. One of the manifestations of apoptosis is the translocation of phosphatidylserine (PS) from the inner membrane to the outer side of the plasma membrane. Externalization of PS was studied by an Annexin V binding assay. Briefly, cells were washed with PBS twice and re-suspended in binding buffer containing 0.01 M HEPES, 0.14 mM NaCl and 2.5 mM CaCl₂. A cell suspension (1x10⁵ cells in 100 μl) in the binding buffer was incubated with 5 μl of FITC-labeled Annexin V (BD PharMingen, San Diego, CA) and vital dye propidium iodide for 15 min in the dark at room temperature. After incubation, the fluorescence of the cells in 100 μl of binding buffer was measured simultaneously in a BD FACS Calibur and analyzed with the operating software of the instruments (CellQuest; BD PharMingen). Data acquisition and analysis were done with the CellQuest and WinMDI programs.

cAMP levels. Following centrifugation of cultures, the supernatants were removed and 0.1 N HCl with DMEM-F12 medium (1:1) was added to the cells to stop the reaction at each 24-h interval until 96 h. Briefly, 25 μl of the samples was used to measure cAMP levels. cAMP accumulation was measured in the supernatants, as described previously with some modifications (28). cAMP was determined by radioimmunoassay using the acetylation protocol (29). High-affinity rabbit anti-cAMP antibody was raised in our laboratory by using BSA-conjugated CAMP. Succinyl-cAMP tyrosine methylster (ScAMP-TME) was iodinated by the chloramine-T method (30). Mono- and di-iodo ScAMP-TME were used as tracer ligands for radioimmunoassay and they were then purified by gel-filtration chromatography (Sephadex G 25 superfine), equilibrated and eluted with 1 M sodium acetate (pH 5.0). Results are given as pmol/ml.

Transmission electron microscopy. Harvested spheroids were fixed with 2.5% glutaraldehyde in 0.1 M sodium cacodylate buffer and post-fixed in 1% osmium tetroxide in 0.1 M sodium cacodylate buffer for 1 h at 4°C. Cells were incubated in 1% uranyl acetate for 1 h at 4°C, dehydrated in a graded acetone series and embedded in Epon 812. Samples were cut on a microtome (Leica MR 2145, Heerbrugg, Switzerland) and 70-nm thick sections were mounted on copper grids. They were stained with 5% uranyl acetate and counterstained with Reynold's lead citrate. Sections were examined with a Jeol-Jem 1011 transmission electron microscope.

Scanning electron microscopy. C6 glioma cells were seeded on microslides in 24-well cultures plates containing 1 ml DMEM-F12 medium at a concentration of 5x10⁴ cells/well. After attachment of the cells, drugs in a concentration of 10 μM alone and in combination, were applied to cells. Cells were fixed with 2.5% glutaraldehyde in 0.1 M sodium cacodylate buffer (pH 7.4) for 1 h at 4°C at the 24 and 72 h. Cells were washed twice for 10 min with 0.1 M sodium cacodylate buffer and post-fixed in 1% osmium tetroxide for 1 h at 4°C. Cells were dehydrated in graded acetone series and incubated in amyl acetate. Microslides were critical-point dried, sputter coated with gold-palladium and observed by SEM (Jeol-JSM-5200; Jeol, Tokyo, Japan) Photographs were taken at 25 kV using various magnifications and angles.

Statistical analysis and determination of synergism. SPSS 12.0 statistical software (SPSS, Inc., Chicago, IL, USA) was used for the statistical analysis. The statistical comparison of all results, except for cAMP levels, were analyzed by using the Student’s t-test. Alterations in cAMP levels were analyzed by one-way ANOVA test with post hoc analysis using Tukey HSD at the 24 h and the Welch test with Dunnett T3 test at the 72 h. Data are represented as the mean ± SD, as appropriate, for at least three separate experiments. Results were considered significant with p<0.0005, p<0.001, p<0.01 and p<0.05.

Synergy was determined as described previously (31-33). By this method, synergy was determined using the following equation: CI=(Dₓ/ (Dₓ)+Dₓ) for CIMP and the combination group was analyzed by the following equation: CI=(Dₓ/ (Dₓ)+Dₓ) for the combination of treatment with IM and CIMP. They were calculated for each cell treatment with IM and CIMP alone and in combination, respectively. The combination of IM and CIMP showed a synergistic effect with a CI value of 1 indicated an additive effect, a CI value >1 indicated a synergistic effect and a CI value <1 suggested an antagonistic effect.

Results

Cell proliferation. The effects of single and combination drugs on cell proliferation of C6 glioma is shown in Fig. 1A. The cell number of the control group showed a proportional increase as a function of incubation time up to 96 h. In contrast to the control group, all drug treatments reduced the number of cells at all times throughout different rates. Treatment of cells with IM and CIMP in combination resulted in a potent reduction in the number of cells at 72 h by 54 and 87%, respectively (p<0.001 and p<0.0005, respectively). The combination of IM and CIMP caused a 45, 81, 89 and 87% reduction for different times of incubation (p<0.05, p<0.01, p<0.001, respectively). The reduction values of combination of IM and CIMP caused a 45, 81, 89 and 87% reduction for different times of incubation (p<0.05, p<0.01, p<0.001, respectively). The reduction values of IM and CIMP were different from each other after the first 24 h (p<0.01 and p<0.001, respectively). In contrast, the reduction in the number of cells following CIMP treatment and the combination group was different from each other only at 48 h following treatment (p<0.05). The inhibition by the combination group on cell proliferation was synergistic at 96 h.

Cell viability. The cell viability of the group was 90% at 24 h and up to 99% at 96 h. Both single and combination treatments reduced the cell viability through 48 h; the most effective in reducing cell viability was the combination group (13%, p<0.05). After 48 h, all drug treatments lost their
inhibitory effect on cell viability and the percentage of trypan blue-positive cells were the same and/or similar to the control group and to each other (97, 97 and 96% at the 96 h for the IM, CIMP and combination groups, respectively; Fig. 1B).

**Cell cycle kinetics**

**DNA synthesis (S-phase) analysis of monolayer cultures.** Cells with red-stained nuclei were considered BrDU-positive. As shown in Fig. 1C, C6 glioma cell nuclei were positive for BrDU in the control group with BrDU-LI values of 37.53% at 24 h and 43.26% at 96 h. Treatment of cells with IM and CIMP resulted in a potent reduction of BrDU-LI values at 48 and 96 h, with values of 69 and 79%, respectively (p<0.05). Combining IM with CIMP did not have much effect on the BrDU-LI throughout the 96 h of culture and the potent inhibition of the BrDU-LI value (87%) persisted at 24 and 48 h (p<0.05). The decrease of BrDU-LI by the combination represented a synergistic effect at 24 and 72 h.

**DNA synthesis (S-phase) analysis for spheroid cultures.** Twenty-five spheroids from all experimental groups, including the control group, were evaluated for 24 h. Cells with red-stained nuclei were considered BrDU-positive. The control group had a healthy pattern of S-phase fraction and many of the C6 glioma cell nuclei were observed to be positive for BrDU with a BrDU-LI value of 8.7% (Fig. 2A). In comparison to controls, IM and CIMP alone reduced the BrDU-LI values to 3.72 and 2.43%, respectively. In addition, light microscopy photographs supported this statistical values that the spheroids of IM and CIMP groups showed less BrDU-LI positive C6 glioma cells (Fig. 2B and C). When the combination group was compared with the control group, a decrease in the BrDU-LI value to 1.96% was observed. In comparison to the other drug groups, the combination group seemed to have no significant difference as the BrDU-LI value was very similar to the CIMP group (Fig. 2D). In comparison to the controls, either drug treatment alone or in combination was not statistically significant.

**Annexin V-FITC and PI double staining.** Fig. 3 clearly shows different levels of apoptosis induced by the two agents when used singly. The percentage of apoptotic cells in the IM group (14.19%) was modestly increased when compared with the control group (9.82%) at 24 h, but at 96 h treatment with IM decreased the percentage of the apoptotic cells to 9.09%. In comparing the control group and the IM group, CIMP potently induced apoptosis i.e., 35.49 and 80.49% at 24 and 96 h, respectively. However, when IM with CIMP were used, the level in combination of apoptosis was enhanced and greater than additive (61.95%) at 24 h. The percentage of apoptotic cells induced by the combination group (86.52%) were similar to the CIMP group (80.49%) at 96 h.

**cAMP levels.** The cAMP levels of the control group showed a proportional increase from 8.20 pmol/ml at time 0 to 37.47.
pmol/ml at 24 h and 61.61 pmol/ml at the 72 h. cAMP levels of the IM group (37.68 pmol/ml) were almost the same as the control group at 24 h, but IM inhibited cAMP levels similar to the control (31.26 pmol/ml) at 72 h (p<0.05). In comparison to the control and IM groups, CIMP potently decreased cAMP levels to 19.14 pmol/ml at 24 h (p<0.05) and 2.68 pmol/ml at 72 h (p<0.01). The combination of IM with CIMP mildly decreased the levels to 30.40 pmol/ml at 24 h (p>0.05), but this group potently decreased cAMP levels to 5.19 pmol/ml at 72 h (p<0.01). The most potent inhibitory effect among all drug treatments and time intervals was observed in the CIMP group (Fig. 4).

Morphology. We discriminated the ultrastructural differences between the control group and the drug-applied groups by using TEM and SEM.

Spheroid morphology by TEM
24 h of culture. The control group exhibited normal morphology, characterized by fine texture of nuclear chromatin defined as oval, heterochromatic nuclei, intact nuclear membrane, a cytoplasm rich in secretory materials, proportional organelles distributed in the cytoplasm, tubular structure of mitochondria, intact cytoplasmic membrane, and many microvilli, which were in contact with other cells (Fig. 5A). The IM-treated group exhibited a smoothing of the cell surface, characterized with decreased, short microvilli, but intact nuclear membrane, pyknotic nuclei, alterations in mitochondria, as delineated by cristae damage and mitochondrial swelling, packaging of small vacuoles, and presumably autophagic vacuoles with lamellar bodies inside, and huge vacuoles, which may be the result of small vacuole fusion. Vacuoles were localized at the peripheral region of the spheroid (Fig. 5B). Mitochondria which possessed outer membrane damage and disrupted or lost cristae were observed in the CIMP group. In addition, presumably autophagic vacuoles, which were in different dimensions with lamellar bodies inside were also seen in this group, similar to the IM group (Fig. 5C). The extension of the nucleus to the cytoplasm, like a pseudopod, was one of the notable finding observed. Apoptotic bodies were also observed by the effect of CIMP. The contact loss of the cells localized at the proliferation zone was the other effect of the drug. The combination of IM with CIMP reflected the total effects of the two drugs displayed when they were administered alone.
Some morphological changes of apoptosis were observed as nuclear fragmentation, increased autophagic vacuoles, mitochondrial damage with disrupted outer membrane and cristae loss and a smooth cell surface (Fig. 5D).

96 h of culture. The presence of a few vacuoles was the only different morphologic alteration observed at 24 h confined to the control group. The presence of zonula adherens (presumably), endoplasmic reticulum and ribosomes was also noteworthy (Fig. 6A). There were no morphological changes observed in the endoplasmic reticulum regions and cell contact with adjacent cells in the IM-treated group. The same morphological changes were observed as at 24 h, but the degree of mitochondria damage such as cristae loss, membrane loss, and the number of vacuoles (foamy vacuolated cytoplasm), including autophagic vacuoles, was increased in this group at 96 h (Fig. 6B). Vacuole fusions were also observed. The observed morphologic alterations differed from those at 24 h and consisted of the increase of autophagic vacuoles in the CIMP group. Separation of the cells localized at the proliferation zone was also observed (Fig. 6C). The combination of IM with CIMP treatment exhibited morphological changes as cell contact loss at the proliferative zone of spheroids, invagination of the nuclear membrane, and the markedly increase in autophagic vacuoles (Fig. 6D). The presence of big lamellar bodies in the nucleoplasm and plasma membrane fusion was the most interesting features which were observed at that time (Fig. 7A and B). The small components of lytic cells which could be a result of apoptosis were located at the outer part of the spheroid (data not shown).

Cell morphology by SEM

24 h of culture. The control group of C6 glioma cells had a spindle-shaped appearance and frequent, active mitotic division phase was observed under SEM. In the IM group, cells changed from a spindle-shaped appearance morphology to random formation to an angular morphology. IM led to a decrease in the number of microvilli. Some cells in the IM group possessed cell surface knobs characterized by a protruding surface. In the CIMP group, some cells possessed an angular morphology and some cells changed to a globular appearance like a ball, and CIMP led to a sharp decrease in the number of microvilli and cell surface extensions. In the combination group, three different cell types could be observed as follows: i) a wide body with an angular morphology with short cell surface extensions; ii) a globular appearance like a ball; and iii) cell surface knobs characterized by a protruding surface. Apoptotic bodies were prominent in this group. The presence of three different cell types which
Figure 6. Transmission electron microscopy (TEM) view of C6 glioma spheroids at 96 h. (A) Control group (original magnification, x15000). (B) Imatinib group (original magnification, x6000). (C) Chlorimipramine group (original magnification, x12000). Wavy arrow represents the separation point of cell from the proliferation zone of spheroid. (D) Imatinib with chlorimipramine (original magnification, x5000). av, autophagic vacuole; er, endoplasmic reticulum; n, nucleus; mi, mitochondria; mv, microvillus; v, vacuole; vp, package of vacuoles.

Figure 7. Transmission electron microscopy (TEM) view of C6 glioma spheroids at 96 h. Imatinib with chlorimipramine (original magnification, x4000). (A) One arrow represents the plasma membrane fusion. Wavy arrows represent presumably the apoptotic bodies. (B) The presence of big lamellar body in the nucleoplasm (original magnification, x5000). av, autophagic vacuole; cr, remnants of cell; er, endoplasmic reticulum; lb, lamellar body; n, nucleus; nu, nucleolus; mi, mitochondria; mv, microvillus; v, vacuole; vp, package of vacuoles.
was observed by the combination group represented the enhanced cytotoxicity of IM.

96 h of culture. Healthy C6 glioma cells which were in a spindle-shaped appearance were observed in the control group (Fig. 8A). In the IM group, many cells possessed an angular morphology and IM led to a sharp decrease in the number of microvilli and cell surface knobs, which caused an appearance like a protruding surface (Fig. 8B). In the CIMP group, many cells changed to possess a globular appearance like a ball with short cell surface knobs characterized by a protruding surface (Fig. 8C). In the combination group, three different cell types were observed as follows: i) a wide body with an angular morphology with short cell surface extensions; ii) a globular appearance like a ball with short extensions; and iii) no specific appearance with cell surface extensions or with a protruding surface (Fig. 8D).

Discussion

The treatment of malignant brain tumor glioblastoma multiforme with effective chemotherapeutic regimens remains a challenge. One of the most promising developments in chemotherapy has been to substitute cytotoxic agents that inhibit non-specific targets in cellular pathways that contribute to carcinogenesis and resistance (25,26). It has been reported that IM inhibits the growth of glioma cells in preclinical models, but its poor brain distribution by P-gp limits its efficacy in patients (34). In this study, we investigated whether CIMP, which has a specific effect on mitochondria, could increase the cytotoxic effect of IM.

In the present study, treatment with IM and CIMP in combination was slightly better than single agents using both monolayers and spheroid cultures of malignant C6 glioma cells. The results of the present study did not fully support the hypothesis that CIMP could induce a synergistic effect on IM-induced cytotoxicity. By using a number of different techniques, the effect of CIMP on IM toxicity on C6 glioma cells revealed the following: i) the effect of the combination group on cell proliferation was synergistic at 96 h; ii) the cell viability results were additive only at 48 h; iii) the decrease of BrDU-LI represented a synergistic effect at 24 and 72 h; iv) the decrease in the percentage of spheroid culture cells in S-phase; v) reduced cAMP levels; and vi) the combination group represented a synergistic effect in apoptosis. In addition, for autophagy, the results varied from IM as single agent to IM with CIMP in combination and resulted in two different findings: single IMP treatment led to chemoresistance/cell survival and the combination of IM and CIMP led to apoptosis.

The most interesting findings of the present study was determined by transmission electron micrographs of spheroids were the apoptosis results. The combination treatment showed a significant potentiation of apoptosis as compared to
findings suggested that the two drugs synergize with each other. These results showed that IM increased the fraction of apoptotic cells very slightly and CIMP markedly induced and increased apoptosis at all time intervals. However, the findings with the combination group were somewhat surprising; although a low apoptotic fraction in the IM group and a high apoptotic fraction in the CIMP group were observed, the weak effect of IM on the induction of apoptosis, in accordance with the results of transmission electron micrographs, interest was increased by CIMP in the combination group. The effect of CIMP on the apoptotic index was more potent when it was combined with IM. These findings suggested that the two drugs synergize with each other. In addition to TEM micrographs of spheroids, the scanning electron micrographs of monolayer cultured cells supported the increase in the apoptotic index by the combination group. These findings supported the synergistic effect of CIMP on IM-induced toxicity.

In addition, the alterations in cAMP levels of cells also supported the morphologic findings about CIMP-induced mitochondrial damage. C6 glioma cells possess native functional PDGFRs, proliferate in response to PDGF, whose stimulation leads to a well-expressed cAMP and is the first identified second messenger (35,36). The different receptors, chiefly the G-protein coupled receptors and growth factor receptors are responsible for cAMP accumulation in cells that cause different physiologic outcomes, and changes in cAMP levels effect the selective activation of cAMP-dependent protein kinase-A isoforms. The chief source of cAMP is from ATP, which is generated by mitochondria (37,38). In our study, the effective inhibition of cAMP levels was observed with the CIMP group for all time intervals, but IM showed its inhibitory effect at 72 h. The inhibitory effect of the combination group on cAMP levels also increased at 72 h. The alterations of cAMP levels could also be interpreted as the inhibitory effect of IM on growth factor receptors, such as PDGFR.

Spheroids are known to be in vitro-like tissue-culture representatives and the most adapted model to keep the in vitro resistance properties of cells. Spheroids more closely mimic the real biologic environment by providing cell-to-cell adhesion and signalling. Due to these characteristics, spheroids quite realistically represent the results of the drug effects by including limitations in penetration, distribution, and feedback mechanisms in cell signaling (39,40). In this study, similar to the BrDU-LI results of monolayer cultures, all drug applications reduced the BrDU-LI in comparison to the controls. In comparison to the effects of singly applied drugs, little more decrease in BrDU-LI was observed in the combination group. The effect of CIMP could not be defined as antagonistic, additive, or synergistic, although a strong synergistic effect was observed in transmission electron micrographs of spheroids. The prominent morphologic changes were observed in the spheroid micrographs, but the combination group seemed ineffective in reducing cells in S-phase. Because spheroids are the bridge between in vitro and in vivo, these findings suggest that further studies (i.e., high dose applications) should be done to achieve the effectiveness of these drugs in the spheroid model.

Alterations of spheroid morphology revealed that IM induced and increased the cellular autophagy machinery in our study. Taken together, in the present study, especially in light of findings of electron micrographs with Annexin V-FITC and PI double staining and cAMP results together, the effect of IM could be interpreted in two different ways: i) three different studies regarding the IM effect on autophagy were done by Ertmer et al (41) and Carew et al (42,43). Ertmer et al (41) described, for the first time, the process of autophagy with the mode of action of IM. They considered that autophagy may promote apoptosis in cancer cells. They revealed that the IM dose-dependently activates the cellular autophagy machinery in mammalian cells, independently of tissue type, species origin, or the immortalization status of cells. When IM was used in combination, the induction of autophagy by IM might represent an additional mechanism to induce apoptosis in C6 glioma cells, like the study of Ertmer et al (41). Besides, if the mode of action of CIMP was evaluated, the induction of autophagy by CIMP might represent an additional mechanism of CIMP to induce growth arrest and promote apoptosis in C6 glioma cells and this effect of this drug was prominently seen in the combination group. ii) In contrast to Ertmer et al (41), Carew et al (42,43) considered that autophagy may promote cancer cell survival and play important roles in chemoresistance of cancer to some therapeutic agents, which typically induce an apoptotic response. They showed that disrupting autophagy augmented the anticancer activity of the histone deacetylase (HDAC) inhibitor suberoylanilide hydroxamic acid (SAHA) in IM-refractory chronic myelogenous leukemia (CML) and in IM-resistant primary CML cells from patients bearing mutations in Bcr-Abl, including the T315I mutation that causes resistance to currently utilized tyrosine kinase inhibitors and translates into a very poor clinical prognosis (41,42). When IM is used singly, as shown by Carew et al (42,43) the induction of autophagy in the IM-applied group might represent chemoresistance of C6 glioma cells to this drug.

In contrast to the findings of electron micrographs, Annexin V-FITC and PI double staining, cAMP levels, cell viability, and proliferation (Fig. 1A and B), and the BrDU-LI of spheroids (Fig. 2) did not indicate that the combination is more potent than the single agent alone, with minor exceptions. The findings that showed the efficiency of the combination group vs. single applications could be hypothesized to be related to CIMP's action: CIMP penetrates plasma membranes easily because of its lipophilic structure (44). Xia et al (45) showed that CIMP induced apoptosis in CML HL-60 cells via increased ROS generation, which
was followed by an increase of caspase-3 activity. They proposed that CIMP enhanced the level of ceramide which was involved in ROS production by interacting with complex III of the ETC. Likewise, Xia et al (45) and Levkovitz et al (46) found CIMP to induce apoptosis in C6 glioma caspase-3-like activation and cyt-C-release. Daley et al (47) studied the mode of action of CIMP in primary cultures of glioblastoma taken from human biopsies and found that CIMP killed cells through inhibiting glioma cell oxygen consumption by binding complex III, resulting in caspase-3 activation. In the present study, CIMP could kill C6 glioma cells through a mitochondria-dependent fashion, as indicated in studies above. P-gp is a transmembrane protein that engages in drug efflux in an ATP-dependent fashion. This transports serine or threonine phosphorylation sites which were phosphorylated by several kinases such as PKC- and cAMP-dependent protein kinase (46). IM is a substrate for P-gp (12). CIMP inhibited ATP production via inhibition of mitochondrial respiration and this resulted in the inhibition of cAMP production. In the present study, CIMP was able to facilitate the penetration of IM through the inhibition of P-gp.

Although, CIMP is not considered as a typical chemotherapeutic drug, the present study did not completely substantiate the synergistic cytotoxic effect of IM combined with CIMP on C6 glioma cells and a spheroid model. Briefly, CIMP appears to act with IM only in monolayer cultures of C6 glioma cells, with few exceptions. Further studies should be done to provide the basis for a possible use of IM and CIMP in combination in vivo and to consider CIMP as ‘a promising compound’ for overcoming IM resistance in the treatment of glioblastoma.

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