GL15 and U251 glioblastoma-derived human cell lines are peculiarly susceptible to induction of mitotic death by very low concentrations of okadaic acid

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Abstract. The challenging possibility of selectively inducing mitotic death in tumor cells by combining genotoxic agents with the inhibition of G2 checkpoints of the cell cycle is the subject of intensive investigation. We show that very low concentrations (3.5 and 5 nM) of okadaic acid induce mitotic death in two glioblastoma cell lines, in the absence of genotoxic agents. At the concentrations used, the main target of okadaic acid action is protein phosphatase 2A (PP2A), an enzyme deeply involved in the negative control of cell-cycle progression. The peculiar susceptibility of glioblastoma cells to induction of mitotic death by very low concentrations of okadaic acid must be related to an impairment of PP2A activity and to a specific deficiency in some cell-cycle checkpoints. In addition to its ability to induce abnormal mitoses in actively proliferating glioblastoma cells, okadaic acid possesses the ability to force semi-confluent glioblastoma cells to the M phase of the cell cycle, where they show the same abnormalities observed in actively proliferating glioblastoma cells. In semi-confluent cells the induction of mitotic death involves the activity of both the extracellular signal regulated kinases (ERKs) and the M-phase promoting factor: okadaic acid overstimulates ERK activity, and PD98059 (inhibitor of ERK activation) as well as roscovitine (S)-isomer (specific inhibitor of M-phase promoting factor activity) counteract the induction of mitotic death. Our results show that, without the use of genotoxic agents, it is possible to induce mitotic death in glioblastoma cells by activating the same uncontrolled pathways responsible for the uncontrolled proliferation.

Introduction

Glioblastoma is the most aggressive form of malignant gliomas, brain tumors originating from glial cells or their precursors. In the last few decades, some characteristic genetic alterations have been identified and related to specific biochemical pathways involved in the control of cell-cycle progression. The complex pattern derived from these studies indicates that abnormal expression of growth factors and their receptors, defective cell-cycle checkpoints, and activation of anti-apoptotic activities contribute to the altered proliferative behaviour of glioblastoma cells. In these cells, the G1 checkpoints are deficient and the tumor suppressor, p53, is often inactive (1,2). Despite the large body of information acquired, an effective therapy of human glioblastoma is still lacking. However, it has been reported that, in several human glioblastomas, the highest degree of apoptosis was detected in the areas with the highest mitotic index suggesting that, during mitosis, some glioblastoma cells are susceptible to spontaneous apoptosis (3).

It has been recently suggested that, during mitosis, tumor cells may be more prone to cell death when a combination of deficient cell-cycle checkpoints and DNA damage occurs (4). This suggests the possibility of sensitizing cancer cells to genotoxic agents by the use of small molecules that inhibit the DNA damage-induced G2 arrest (5). Despite the genomic alterations, tumor cells can overcome the G1 check-point controls and reach the G2 phase of the cell cycle where residual checkpoint controls are able to avoid mitotic death (6,7). The G2 checkpoints negatively control the activity of the M-phase promoting factor (the complex p34 cdc2 kinase-cyclin B1), the best characterized effector on G2/M transition (8). Therefore, a pharmacological forced activation of the M-
phase promoting factor could mimic the inhibition of G2-phase checkpoints.

In normal conditions, protein phosphatase 2A (PP2A) can constitute a potential negative control of cell-cycle progression through both the G1 phase and the G2 phase (9). In the G1 phase, the involvement of PP2A could be hypothesized given its role in the downregulation of the Ras/ERK (extracellular signal regulated kinase 1/2) pathway (10-12). In this respect, dephosphorylation of MAPK/ERK kinase (MEK) and ERK, performed by PP2A, could constitute a key mechanism in the inactivation of the ERK pathway (13). In several cellular systems, activation of the ERK pathway is strongly related to the induction of proliferation by transcriptionally activating the numerous mitogen-inducible genes regulated by serum response elements (SREs) (14,15). Moreover, it has been demonstrated that ERKs are able, in late G1, to activate the CDK2/cyclin E complex (16) involved in the control of centriole duplication (17) and, in the G2 phase, to cooperate with Polo-like kinase 1 in the translocation of cyclin B1 (the complex cdc2 kinase-cyclin B1) to the nucleus (18). PP2A has been shown to be deeply involved in the downregulation of the M-phase promoting factor (19-22). Therefore, a forced activation of the M-phase promoting factor could be obtained by PP2A inhibition mimicking an overcoming of some cell-cycle checkpoints.

The most widely used inhibitor of PP2A is okadaic acid (OA), a cell permeable polyether fatty acid produced by marine dinoflagellates (23-26). In fact, although several protein phosphatases are sensitive to the inhibitory action of OA, PP2A is the most sensitive.

In cellular systems, the biological effects of OA depend on the cell type and mostly on the concentrations used: OA can act as a tumor promoter but also as an inducer of apoptosis. The OA concentrations able to induce apoptosis in several cell lines are so high that they do not allow the identification of the specific protein phosphatase and, therefore, the mechanism involved.

In order to specifically inhibit PP2A, we analyzed the effects of very low concentrations (≤5 nM) of OA on GL15 (27-31) and U251 (32) human glioblastoma cell lines. The main effect was the induction of apoptosis occurring during mitosis.

Our results constitute the first demonstration of an apoptotic effect driven by OA at these concentrations and highlight the possibility of identifying glioblastoma-specific biochemical alterations in which the impairment of PP2A is involved. These specific alterations could be the targets for a future pharmacological approach.

Materials and methods

Cell cultures. Cultures of the GL15 glioblastoma multiforme cell line were grown in minimum essential medium (MEM) supplemented with 10% heat-inactivated foetal bovine serum (FBS), 100 IU/ml penicillin G, 100 μg/ml streptomycin, 2 mM glutamine and 1 mM sodium pyruvate. Cultures of the U251 glioblastoma cell line were grown in Dulbecco's modified minimum essential medium (DMEM) supplemented with 10% heat-inactivated FBS, 100 IU/ml penicillin G, 100 μg/ml streptomycin and 2 mM glutamine. The flasks were incubated at 37°C in a 5% CO₂ humidified atmosphere. The medium was changed twice weekly and the cells were subcultivated when confluent.

GL15 and U251 cells were treated with 3.5 and 5 nM okadaic acid (Calbiochem) in culture medium, at the indicated times. The other treatments were performed in the same conditions as the okadaic acid treatment. When the combined action of 25 μM PD98059 (Calbiochem) with okadaic acid was analysed, the compound was added to the culture medium 30 min after the inhibitor. The final concentrations of roscovitine (S)-isomer (Calbiochem) and SB203580 (Calbiochem) were 20 μM and 10 μM respectively. In all experiments, the final concentration of dimethyl sulphoxide (DMSO) was 0.15%. Controls containing the same amount of DMSO were included at each experimental time point.

Morphological analysis of the cells was performed with a contrast phase Nikon microscope. Cell count was performed after trypsinization using a Burker camera.

DAPI staining. Cells were extensively washed with phosphate-buffered saline (PBS) and fixed with 4% paraformaldehyde for 15 min. After washing with PBS, cells were incubated with 2 μg/ml 4,6-diamidino-2-phenylindole (DAPI, Sigma) for 5 min and dried in air. The preparations were observed using a DMRB Leika microscope.

Indirect immunofluorescence. Cells were extensively washed with PBS, immersed in cold methanol, kept at -20°C for 7 min and permeabilized by treatment with PBS containing 0.1% Triton X-100 for 5 min. The cells were then incubated for 60 min at room temperature with a mouse anti-β-tubulin antibody (Sigma), diluted 1:50 in PBS containing 0.3% bovine serum albumin, and washed 3 times with PBS containing 0.1% Tween-20 and twice with PBS alone. After treatment with fluorescein isothiocyanate (FITC, Sigma) conjugated goat anti-mouse (IgG diluted 1:50 in PBS containing 0.3% albumin) and washing 3 times with PBS containing 0.1% Tween-20 and twice with PBS, preparations were incubated with 2 μg/ml DAPI for 5 min and dried in air. The preparations were observed using a DMRB Leika microscope.

ELISA detection of cytoplasmic nucleosomes. Determination of cytoplasmic histone-associated DNA fragments was performed using the Cell death detection ELISA (PLUS) kit (Roche), following the instructions of the manufacturer. The results are expressed as percentage of optical density, resulting from the activity of peroxidase-conjugated anti-DNA-antibody complexed with cytoplasmic nucleosomes of treated cells, compared to the control.

SDS-polyacrylamide gel electrophoresis and immunoblotting. GL15 cell cultures were washed with PBS and scraped with 62.5 mM Tris-HCl (pH 6.8), 2 mM ethylenediaminetetraacetic acid (EDTA), 0.5% Triton X-100, phosphatase inhibitor cocktail II (Sigma), protease inhibitor cocktail (Sigma), 0.1% SDS. The proteins were separated by SDS-PAGE in 10% acrylamide gel by using the Laemmli method (33) and then transferred to nitrocellulose filters according to Towbin et al (34). Immunolabeling of phosphorylated and total-ERKs was
performed following the instructions of the manufacturer. The primary antibodies were a mouse anti-phospho-p44/42 MAPK (Thr202/Tyr204) E10 monoclonal antibody and a rabbit anti-p44/42 MAPK antibody (Cell Signaling Technologies) respectively. The secondary antibodies used to detect phospho-and total-ERK content were the peroxidase-conjugated goat anti-mouse immunoglobulins and the peroxidase-conjugated goat anti-rabbit immunoglobulins (Pierce).

Enhanced chemiluminescence detection was performed using the instructions of ECL™ Western blotting (Cell Signaling Technologies).

**Measurements of cell-cycle by flow cytometry.** After the different treatments, aliquots of the cell suspension were transferred into centrifuge tubes, washed once with PBS (400 x g, 7 min) and successively processed for cell-cycle analysis by propidium iodide (PI)-staining and flow cytometry. Briefly, the cell pellet was resuspended in 0.5 ml of hypotonic fluorochrome solution (50 µg/ml PI in 0.1% sodium citrate plus 0.1% Triton X-100) in 12x75-mm polypropylene tubes (Becton-Dickinson, Lincoln Park, NJ). The tubes were kept at 4°C for at least 30 min before flow cytometric analysis. The PI fluorescence of individual nuclei was measured using a FACScan flow cytometer (Becton-Dickinson, Mountain View, CA) at a wavelength of 488 nm. The percentages of the cells in G0/G1, S and G2/M phases were calculated using CellFIT CellCycle Analysis Version 2.0.2. software.

**Results**

In the GL15 glioblastoma-derived human cell line, the expression of large T and thus of small T antigen of SV40 has been previously demonstrated (35). Therefore, in this cell line the tumor suppressors, Rb and p53, are inactivated by the large T antigen (36) and P2A is partially inactivated by the small T antigen of SV40 (37). In this respect, the biochemical abnormalities derived from SV40 large T antigen expression mirror the behaviour of some human glioblastomas (38).

The effects of 5 nM OA were analysed using several experimental approaches: inspection of cell and nuclear morphology, immunocytochemistry, cell-cycle analysis, and evaluation of cytoplasmic nucleosomes.

Two days after treatment, the analysis of the effects of OA (5 nM) on GL15 cell morphology indicated a strong phenotypical regression characterized by round or polygonal shape and by the absence of process bearing cells, and an induction of cell death characterized by a significant amount of cells which round up and detach from the substrate (data not shown).

The cell count confirmed the induction of cell death: a significant decrease in cell number was observed when the treatment was performed on actively proliferating (1 day of subculturing) as well as on semi-confluent (4 days of subculturing) GL15 cells. The decrease in cell number was 42±13.8% SD in actively proliferating and 32±10% SD in semi-confluent GL15 cells.

When analysing the effects of OA on nuclear morphology by DAPI staining, the relevant result was the presence of abnormal mitoses, characterized by a scattered distribution of chromosomes, apparent either in metaphase or in anaphase. Seldom were these abnormal mitoses observed in control cells. In some cases, there was clear evidence that these abnormal mitoses evolved into apoptosis (Fig. 1A.a, b and B.a, b).

These results seem to indicate that OA induces a kind of cell death which occurs during mitosis. This kind of cell death is sometimes defined as mitotic catastrophe, and it has been recently demonstrated to constitute a special case of apoptosis (39).

Due to the fact that abnormal mitotic spindles may lead to mitotic catastrophe, the spindle morphology was visualized by immunocytochemistry, performed using an anti-ß-tubulin antibody. After treatment with OA, abnormal multipolar or monopolar spindles were observed. In analogy with the results obtained by DAPI staining, abnormal mitotic spindles were seldom observed in the controls (Fig. 1A.c and d).

OA induced an evident increase in the number of mitotic cells (mostly abnormal). In order to quantify this increase, after DAPI staining, we compared the abnormal mitoses in metaphase and anaphase, induced by OA, with normal mitoses in metaphase and anaphase in control cells. The increase was evident when the treatment was performed on actively proliferating and semi-confluent GL15 cells (Fig. 1A.e and B.c). It is interesting to note that the increase is higher in semi-confluent GL15 cells.

In order to get a deeper comprehension of OA effects on GL15 cells, cell-cycle analysis was performed after propidium iodide staining. Two days after treatment, 5 nM OA significantly affected the cell-cycle profiles of both actively proliferating and semi-confluent GL15 cells.

When the treatment was performed on actively proliferating GL15 cells, a significant decrease of the G0/G1 phase of the cell cycle, combined with a significant increase of the sub-G0/G1 peak, was observed. The other phases of the cell cycle seemed to be slightly affected (Fig. 1A.f). Therefore, the main effect appeared to be an induction of apoptosis. It is interesting to note that abnormal mitoses were not detected in the cell-cycle profile as an increase of the percentage of cells in the G2/M phase. The DNA content per cell in the G2/M phase could have been affected by the simultaneous degradation induced by apoptosis. Another possibility is that the increase of the percentage of cells in the M phase was shielded by the simultaneous decrease in the percentage of cells in the G2 phase of the cell cycle.

When the treatment was performed on semi-confluent GL15 cells, a significant decrease in the G0/G1 phase, combined with a significant increase in the S and G2/M phases of the cell cycle, was observed (Fig. 1B.d). Therefore, the main effect appeared to be mitotic arrest.

These results could indicate that the apoptotic event shields the mitotic arrest in actively proliferating GL15 cells, while the mitotic arrest prevails in semi-confluent GL15 cells; however, the DAPI staining indicated that the abnormal mitoses, induced by OA, evolved into apoptosis when the treatment was performed on actively proliferating and semi-confluent GL15 cells. Therefore, if the apoptosis evolves from abnormal mitoses, the apoptotic events could be scattered along the whole cell-cycle profile.

In order to confirm this hypothesis, quantitative analysis of cytoplasmic nucleosomes was performed using the ELISA method: our results indicate that, at concentrations of 3.5 and
5 nM, OA induced a significant increase of cytoplasmic nucleosomes on actively proliferating and semi-confluent, GL15 cells (Fig. 1A and B). This result constitutes a demonstration of the apoptotic effect of OA on GL15 cells.

If the apoptosis induced by OA occurs during mitosis, semi-confluent GL15 cells must be less sensitive to OA action. Semi-confluent GL15 cells, instead, are as susceptible as actively proliferating GL15 cells to the induction of mitotic death induced by OA. The very high percentage of mitotic figures induced by OA in semi-confluent GL15 cells seems to underscore an underlying mitotic stimulus.

Owing to the fact that PP2A is the main target of 5 nM OA, an overactivation of the ERKs and of the M-phase promoting factor could result from PP2A inhibition. Therefore, abnormal activities of ERK1/2 and/or of the M-phase promoting factor were analysed in semi-confluent GL15 cells as possible effectors of OA apoptotic action. Both these activities could be responsible for the hypothesised mitotic stimulus.

Six hours after treatment, OA induced a significant increase in the activity of ERK1/2, as evaluated by using a specific antibody against the double phosphorylated (Thr202/Tyr204) ERK1/2 and normalizing with the anti-total-ERK1/2 antibody. This result does not completely identify the increase in ERK1/2 activity as being responsible for the OA apoptotic effect on GL15 cells.

In order to relate the increased ERK activity to the apoptotic effect of OA, experiments with an inhibitor of ERK pathway activation, PD98059 (40), were performed. The increase in

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**Figure 1.** Effects of 5 nM okadaic acid, 2 days after treatment, on GL15 cells. (A) Effects on actively proliferating GL15 cells: a, DAPI staining, control; b, DAPI staining, okadaic acid (original magnification x600); c, immunofluorescence staining of β-tubulin, control; d, immunofluorescence staining of β-tubulin, okadaic acid (original magnification x400); e, percentage of mitotic cells, as evaluated by DAPI staining (C, control; OA, 5 nM okadaic acid); f, analysis of cell cycle evaluated by PI staining and flow cytometry (C, control; OA, 5 nM okadaic acid); g, ELISA detection of cytoplasmic nucleosomes (C, control; OA 3.5, 3.5 nM okadaic acid; OA 5, 5 nM okadaic acid). (B) Effects on semi-confluent GL15 cells: a, DAPI staining, control; b, DAPI staining, okadaic acid (original magnification x600); c, percentage of mitotic cells, as evaluated by DAPI staining (C, control; OA, 5 nM okadaic acid); d, analysis of cell cycle evaluated by PI staining and flow cytometry (C, control; OA, 5 nM okadaic acid); e, ELISA detection of cytoplasmic nucleosomes (C, control; OA 3.5, 3.5 nM okadaic acid; OA 5, 5 nM okadaic acid). *P<0.05, **P<0.01, ***P<0.001, Student’s t-test.
ERK phosphorylation, induced by OA, was significantly reduced by PD98059 (Fig. 2A). Along with the inhibition of ERK activity, PD98059 was also able to decrease the number of abnormal mitoses, as evaluated by DAPI staining (data not shown), and to partially counteract the apoptotic effect of OA, as evaluated by the quantitative analysis of cytoplasmic nucleosomes (Fig. 2B). The cell number was comparable when the treatment was performed with OA alone or with OA + PD98059.

Due to the fact that the action of PP2A in the down-regulation of the M-phase promoting factor is well documented, the possible involvement of a forced activation of the M-phase promoting factor as an inducer of apoptosis was investigated by using a specific inhibitor of p34 cdc2 kinase, roscovitine (S)-isomer (41).

Roscovitine (S)-isomer partially counteracts the morphological effects of OA on semi-confluent GL15 cells; moreover, when analysing the effects of roscovitine (S)-isomer by DAPI staining, a significant decrease in abnormal mitoses, induced by OA, was observed (Fig. 2C). It is interesting to note that, when the treatment was performed with the roscovitine (S)-isomer alone, the result was the total absence of mitotic nuclear morphologies. This finding confirms the inhibition, in GL15 cells, of p34 cdc2 kinase activity because the phosphorylation of nuclear proteins, performed by cdc2 kinase, is involved in the chromatin condensation at the first stages of mitosis (42).

Roscovitine (S)-isomer significantly counteracted the induction of apoptosis by OA, as evaluated by the ELISA method aimed at quantifying the cytoplasmic nucleosomes (Fig. 2D). The cell number was comparable when the treatment was performed with OA alone or with OA + roscovitine (S)-isomer.

Due to the fact that an apoptotic action induced by these concentrations of OA has never been reported, we hypothesised that the peculiar sensitivity of GL15 cells was related to the partial inhibition of PP2A by the SV40 small T antigen, but we could not rule out the possibility that this behaviour was related to the peculiar genetic and biochemical alterations typical of glioblastoma cells (1,43-45). Therefore, we analyzed the effects of OA on another human glioblastoma cell line, U251 where, to our knowledge, the expression of large T antigen of SV40 has never been reported.

Our results show that 3.5 and 5 nM OA was able to induce mitotic arrest and apoptosis, combined with a strong reduction in cell number, in U251 cells. Two days after treatment with 5 nM OA, the DAPI staining showed the presence of abnormal mitoses (Fig. 3A) and a significant increase in the number of mitotic nuclei in metaphase and anaphase, compared to the control (Fig. 3B). Moreover, 2 days after treatment with 3.5 and 5 nM OA, the ELISA method aimed at quantifying the presence of cytoplasmic nucleosomes showed the induction of apoptosis (Fig. 3C). Therefore, the apoptotic response to these concentrations of OA seems to be an intrinsic property of these glioblastoma cell lines.

Discussion

In this study, we demonstrate that two glioblastoma derived human cell lines are peculiarly susceptible to induction of apoptosis by very low concentrations (3.5 and 5 nM) of OA.
factor maturation. The possibility of selectively inhibiting and PP2A converge on the inhibition of the M-phase promoting factor through different biochemical pathways, some G2 checkpoints approach arose from the fact that, in the G2 phase, although some checkpoint controls. The sensitivity of tumor cells to genotoxic agents is increased by a further inhibition of some G2 checkpoint proteins, including p53, 14-3-3, Chk1 and Chk2.

Although other tumor cell lines, namely Ras-transformed tumor cell lines, have been demonstrated to be particularly susceptible to induction of apoptosis by 20 nM OA (46), the sensitivity of GL15 and U251 glioblastoma cell lines seems exceptional. In principle, while 20 nM OA inhibits both PP2A and PP1A, 3.5 and 5 nM OA mainly inhibit PP2A.

Analysis of our results shows that, in both cell lines, the main OA effects are the induction of abnormal mitoses, characterized by a scattered distribution of chromosomes, and the induction of apoptosis. OA seems to induce a kind of cell death which occurs during mitosis. This kind of cell death is sometimes defined as mitotic catastrophe, and it has been demonstrated that tumor cells are more prone to this kind of cell death induced by genotoxic agents, probably because of the uncontrolled G1 progression and the lack of some G2 checkpoint controls. The sensitivity of tumor cells to genotoxic agents is increased by a further inhibition of some G2 checkpoint proteins, including p53, 14-3-3, Chk1 and Chk2.

Instead of using genotoxic agents, we used low concentrations of OA in order to potentially overcome some cell-cycle checkpoints by specific inhibition of PP2A. This experimental approach arose from the fact that, in the G2 phase, although through different biochemical pathways, some G2 checkpoints and PP2A converge on the inhibition of the M-phase promoting factor maturation. The possibility of selectively inhibiting PP2A in cellular systems is usually prevented because the expression of PP2A is very high when compared with that of other protein phosphatases (9). It is possible that the peculiar response of GL15 and U251 cells to OA was due to an impairment of PP2A activity in glioblastoma cells.

In several human glioblastoma cells, an impairment of PP2A activity could be due either to chromosomal aberrations, involving the regions in which the genes encoding the PP2A Aα or Aβ subunits map (47,48), or to a partial inhibition of PP2A by SV40 small T antigen (37). This impairment of PP2A has been related to the abnormal growth of glioblastoma cells.

PP2A has been proposed to act as a tumor suppressor on the basis of the effects of OA (49), originally identified as a tumor promoter in epithelial cells. Moreover, the involvement of PP2A in control of the Wnt pathway, leading to growth stimulation, has been demonstrated (50).

A strong inhibition of PP2A could be achieved when this enzyme is partially inactive, and our results point out the possibility that this achieved strong inhibition could be responsible for cell death in glioblastoma cells, where some feedback control mechanisms are deficient. In GL15 and U251 cells, along with the uncontrolled G1 progression, the inactivity of the tumor suppressor p53, has been demonstrated (35,51).

We have shown that, in semi-confluent GL15 cells, the activities of the ERKs and of the M-phase promoting factor are involved in the induction of mitotic death by OA.

The best known effects of ERK activation concern the control of the G1 phase of the cell cycle, and we cannot rule out the possibility that the induction of cell death by OA involves an uncontrolled stimulation of the ERK pathway in this phase of the cell cycle. In semi-confluent GL15 cells, the very high increase of mitotic figures induced by OA seems to indicate some kind of mitotic stimulus, possibly affecting the G1 or G2 phase of the cell cycle. Moreover, the involvement of ERKs in the centriole duplication and in the translocation of the M-phase promoting factor to the nucleus has been demonstrated. Therefore, the over-activity of ERKs can contribute to the abnormal maturation of the M-phase promoting factor.

Roscovitine (S)-isomer, the specific inhibitor of M-phase promoting factor activity, significantly inhibited the induction of abnormal mitoses and of apoptosis induced by OA. This result has two main explanations: a) the activity of the M-phase promoting factor is essential for mitotic catastrophe because its inactivity prevents the G2/M transition; b) the inhibition of PP2A promotes an uncontrolled activation of the M-phase promoting factor, mimicking the inhibition of some G2 checkpoints.

Some reports define the action of OA as mitotic arrest, probably because some OA-sensitive phosphatases have been shown to be involved in the exit from mitosis (52); along with the fact that high OA concentrations did not allow the identification of the specific protein phosphatase involved, it is difficult to explain the observed abnormal mitoses characterised by multipolar spindles as simple mitotic arrest. In fact, centriole duplication does not occur at the end of mitosis. It is interesting to note that ERKs and Polo-like kinase 2 have been shown to control centriole duplication (16,53). Moreover, the cooperation between ERKs and Polo-like kinase 1 has...
been reported (18). In glioblastomas, the overexpression of Polo-like kinase 1 (54) and an abnormal activity of the ERKs (55), combined with altered cell-cycle checkpoints, could be involved in the propensity for aneuploidy. The orchestrated phosphorylation of mitotic substrates, which allows the correct cell-cycle progression, must be altered. A further disengagement between these activities should be the basis for the induction of cell death by OA. In fact, the selective inhibition of PP2A causes an imbalance among the basic mechanisms which control the progression of the cell cycle; this imbalance is counteracted in normal cells by a plethora of pathways, but in glioblastoma cells some of these pathways are lacking.

The identification of these pathways will indicate very specific targets for pharmacological approaches, considering the possibility of inducing apoptosis in tumor cells by overstimulation of the same uncontrolled pathways responsible for the uncontrolled proliferation.

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