

Presence of MHC-I in rat glioma cells expressing antisense IGF-I-Receptor RNA

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Abstract

The supposed immunogenic character of glioma cells transfected with antisense IGF-I-Receptor (IGF-I-R) expression vector was tested for the presence of MHC-I currently present in cells of IGF-I antisense type. C6 rat glioma cell line was comparatively transfected in vitro with IGF I antisense (pMT-Anti-IGF I) or IGF I Receptor antisense (pMT-Anti-IGF I R) expression vectors. The wild and transfected cells were examined for the presence of IGF-I and MHC-I molecules. Using RT PCR technique, the transfected "antisense" cells showed total inhibition of IGF-I. The both transfected cultures of IGF-I and of IGF-I -R type were positively stained for MHC-I. Moreover "antisense IGF-I-R" cells as compared to "IGF-I antisense" cells showed slightly higher expression of MHC-I. The transfected cells showed also the feature of apoptosis in 60% of cells. The immunogenicity of IGF-I-R antisense glioma cells is related to MHC-I presence; therefore both approaches of antisense IGF-I and of antisense IGF-I-R could be use in paralel for cellular therapy of glioblastoma.

Key words: glioma, IGF I, IGF-I Receptor, antisense, MHC-I.

Introduction

Different strategies of gene therapy for treatment of gliomas have been proposed comming from 1990s [1, 2, 3]. Our strategy

treatment is based on antisense [4, 5] anti - IGF-I (insulin-like growth factor I) technology [1]. IGF-I is a 70-amino acid polypeptide involved in tissue growth and differentiation [6, 7, 8, 9].

Deregulated expression of growth factors and/or their receptors, and especially of IGF-I, is associated as well with growth as with pathology of different diseases, including tumors [9, 10, 11, 12]. IGF-I acts via specific IGF-I receptor and subsequent activation of a protein tyrosine phosphorylic signal transduction cascade, similar to that of insulin action [13, 14]. IGF-I, mediated by IGF-I receptor has been reported to stop the apoptosis pathway in a variety of cell lines [9, 15]. The action of IGF-I on cellular metabolism depends on binding proteins (IGFBP), which prolong the half life of this factor and modify its interaction with receptor [16, 17]. The block of IGF-I synthesis, induces apoptotic and immunogenic phenomenons [18].

IGF-I is expressed at high levels in some nervous system-derived tumours, e.g. glioblastoma [19, 20] and tumour cell lines like C6 and CNS-1 rat gliomas [21, 22, 23].

Rat C-6 glioma cells become down-regulated in production of IGF-I when transfected with vector expressing IGF-I antisense RNA or IGF-I triple-helix forming oligonucleotides. The transfected cells lose tumorigenicity and elicit tumor specific immunity which leads to cure of established tumors. The immune response was shown to be mediated mainly by T lymphocytes as indicated by an abundant infiltration of the CD8 T cell subset in tumor tissues [12, 24]. Here we examine the possibility that the suppression of intracellular IGF-I in rat glioma cells, using antisense IGF-I-Receptor approach, alters the cell phenotype and the production of MHC-I as it was previously observed in IGF-I antisense approach [25, 26], and that this process is also associated with increase in programmed cell death - apoptosis.

Material and methods

Cell culture. C6 rat glioma cell line [21], was used. Cells were cultured in DMEM+F12 (v/v) (GIBCO-BRL) supplemen-

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ted with 10 % FCS, 2mM glutamine, 100 U/ml penicillin and 100 µg/ml streptomycin, at 37° C and 5 % CO₂. Hygromycin B (Boehringer Mannheim) at a concentration of 0,5 mg/ml was added 48 hours after transfection to select for transfected cell line. Then, the concentration of hygromycin B was changed to 2 mg/ml after one week and maintained with each change of fresh medium during the next 3-4 month.

Plasmids

The episome based plasmids pMT-Anti IGF-I and pMT-Anti IGF-I-Receptor were constructed as previously described [1,3,27]. The vector pMT-EP, under the control of the metallothionein, MT-I, inducible promoter was used as its base. The cassette contains the Epstein-Barr Virus origin of replication and the gene encoding nuclear antigen I which together drive extra-chromosomal replication. Down stream of the insertion site is a poly A termination signal followed by the hygromycin B and ampicillin resistance genes. The vector pMT-EP with the lac-Z reporter gene, as insert was used in control experiments [28].

Transfection

Cultures of cells, 60-80 % confluent, were transfected in 6-well plates utilizing a ratio of 1 µg plasmid DNA per 400 000 cells. The FuGENE 6 Transfection Reagent (Boehringer Mannheim) was used according to the supplier's instructions. To determine the efficiency of transfection, the process was carried out using the pMT-EP construct containing lac-Z as a reporter gene. Cell cultures were washed in PBS and incubated at 37° C in the presence of the staining solution which contained 5mM K₃Fe(CN)₆, 2mM MgCl₂, 0,8 mg/ml X-gal made in PBS.

The cell lines, transfected and non transfected were comparatively analyzed for expression of IGF-I, IGF-I-R and MHC-I in the presence and absence of ZnSO₄ (stimulation of IGF-I metallothionin promoter).

RT PCR (Reverse transcriptase - polymerase chain reaction)

RNA from cells was isolated using High Pure RNA Isolation Kit (Roche Diagnostics GmbH nr 1828665), and the obtained mRNA was verified in 1,5% agarose gel stained with ethidium bromide. The applied components of RT PCR were used according Reverse Transcription System Promega Corporation (nr A3500). Samples composed as mentioned in Table 1. were amplified for 30 cycles.

Each cycle consisted of the following steps: denaturation at 94°C for 30 sec, then at 65°C for 30 sec.; annealing for 60 sec, primer extension at 72°C for 5 min in a DNA thermal cycler (GeneAmp PCR System 2700, Applied Biosystem).

The following primers were used for RT PCR study of:

IGF-I:

Primer "a"

Forward: TAG TCC CTG CCT CTT AAG AG

Reverse: AGG GGC GTT AAA ACT TGG GT

(sequence according "rgd" Rat Genome Database; primers synthesized in Institute of Biochemistry and Biophysics, Polish Academy of Sciences, Warsaw).

Table 1.

Components	Sample (µl)
cDNA	2
dNTPs, 10mM	1,8
MgCl ₂ , 25mM	7,5
RT 10x Bufor	9,8
rev IGF-I primer, 20µM	2
dir IGF-I primer, 20µM	2
DNA Pfu Polymerase (2,5)	0,5
Nuclease free water (100µl)	74.4

IGF-I:

Primer "b"

Forward: TCT GCT GAA GCC TAC AAA GT

Reverse: TTC CTC AAG CAG CAA AGG AT

IGF-I-Receptor:

Primer "c"

Forward: GAC AGT GAA TGA GGC TGC AA

Reverse: TCT CCA CCT CTG GCC TTA GA

MHC-I:

Primer "d"

Forward: ACA CTC GCT GCG GTA TTT CT

Reverse: CCT TGG CTT TCT GTG TCT CC

(primers synthesized in Institute of Pharmacology, Polish Academy of Sciences, Cracow)

Amplification products were analyzed in 1,5 % agarose gel; each well was loaded by 0,5 ug DNA stained by ethidium bromide. Standard markers were used (PCR markers, Promega Corporation, nr G316A).

Results

Morphological characterisation of tumoral cells.

C6 glioma cells demonstrated morphological characteristics similar to those previously described [1] (Fig. 1). The percent of IGF-I positive cells ranged from 70-90%. Each cell line was subcloned to obtain IGF-I positive clones. IGF-I antisense transfected cells (pMT-Anti IGF-I) as well as IGF-I-R antisense transfected cells (pMT-Anti IGF-I-R) did not stain immunocytochemically for IGF-I [1, 27]. The frequency of transfection for the pMT-EP- LacZ vector, as determined by staining with X-gal at 24 hours after transfection, was in between 3-10 % for glioma lines (Fig. 2). In cultures of both IGF-I antisense and IGF-I-R transfected cells, approximately 40-50 % of the cells became rounded and detached. These cells were all apoptotic. In contrast, evidence of apoptosis was observed in less than 5-10 % of non-transfected cells. The apoptotic changes occurred within 5-6 hours after incubation of transfected cells in the presence of 60 microM of ZnSO₄. These results were confirmed in six separate experiments (P < 0, 01).

Figure 1. In vitro culture of C6 glioma cells. The cloned cells showing oftenly elongated shape, present a pseudo-fibroblastic character. The apoptosis phenomenon, oval cells with vacuolization is rare (less than 5%). X400

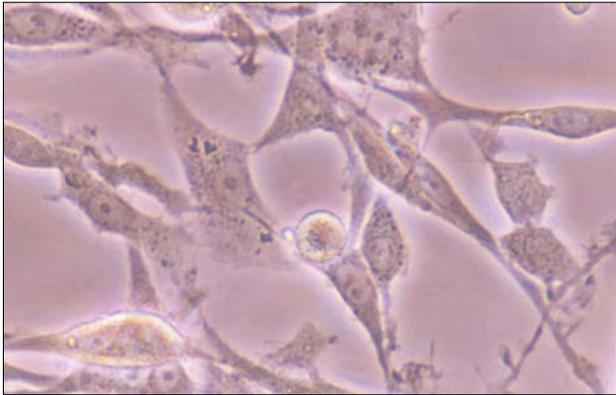
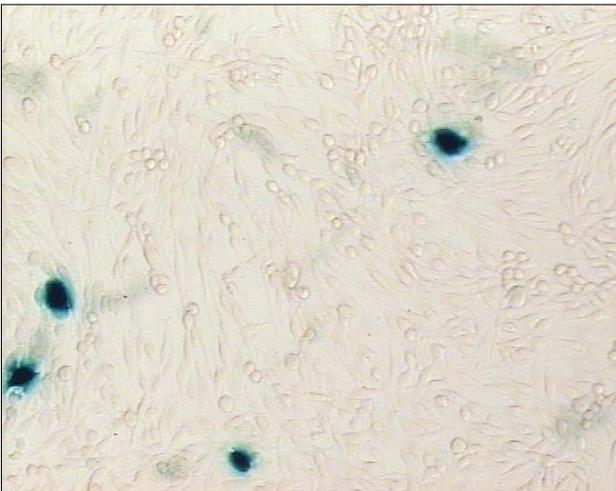


Figure 2. In vitro C6 glioma cells transfected with "antisense" pMT-Anti IGF-I-R vector and with pMT-EP construct containing lac-Z as a reporter gene. The transfected cells are morphologically different from non transfected cells; they are all elongated. Note blue-green cells (X-gal staining) of cells transfected with beta-galactosidase expressing vector (lac-Z). X400



Molecular characterization of tumoral cells.

Analysis of C6 rat glioma, parental, non transfected cells is shown in the RT PCR gels (Figs. 3, 4a and b). Using primer for IGF-I and IGF-I-R, the presense of these two genes was demonstrated in non transfected cells. The DNA of non-transfected cells is distributed in 300-400 bp (Fig. 3) and 200 bp bands (Figs. 4a and b) corresponding to the different primers.

As far as transfected cells are considered, the presence of MHC-I was confirmed in "antisense IGF-I cells" [12] using RT PCR technique. These cells showing absence of IGF-I, demonstrated the expression of MHC-I after stimulation with Zn SO4 (Figs. 5a and b). The same striking phenomenon was observed in "antisense IGF-I-Receptor" cells, also stimulated with ZnSO4 (Figs. 6a and b), indicating the presense of MHC-I. Non stimulated cells show very weak bands of 200 bp or their absence using MHC-I primers.

Figure 3. The example of RNA isolation and of amplified DNA in C6 non transfected, parental cells. M - marker (PCR Markers, Promega Corporation, nr G316A. 1, 2, - isolated samples of RNA. 3, 4, 5, 6, 7 - amplified samples of DNA: note 300-400 bp bands confirm the expression of IGF-I using IGF-I primer "a" (see Material and methods).

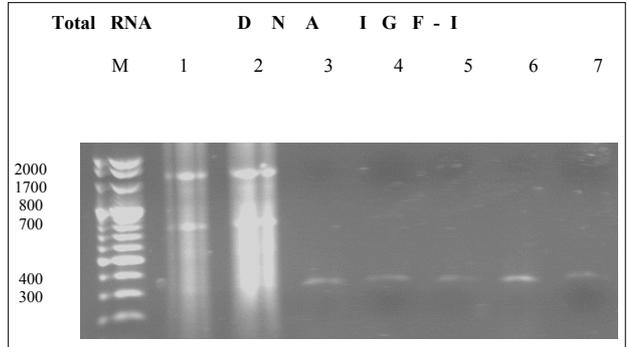


Figure 4a. C6 parental cells. The expression of IGF-I demonstrated by RT PCR. M- marker; 1,2,3 - 200 bp bands of amplified DNA using IGF-I primer "b" (see Material and methods).

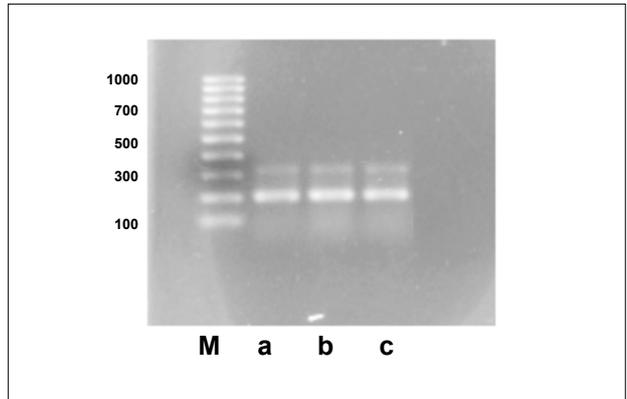
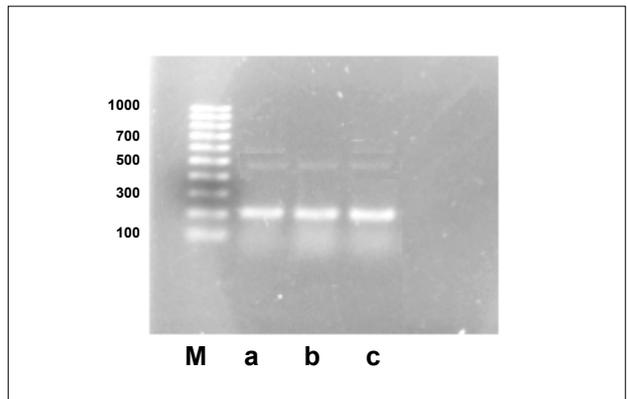


Figure 4b. C6 parental cells. The expression of IGF-I-Receptor demonstrated by RT PCR. M- marker; 1,2,3 - 200 bp bands of amplified DNA using IGF-I-R primer "c"; 300 bp is non specific (see Material and methods).



Discussion

Other studies have already shown, that the antisense strategy directed to the receptor of IGF induces a programmed cell death of tumor cells [9, 18, 22, 29]. As antitumor IGF-I strate-

Figure 5a. C6 glioma cells transfected with pMT-Anti IGF-I. Absence of expression of IGF-I demonstrated by RT PCR. M- marker; a, b, c, d, e, f - amplified DNA using IGF-I primer "b" (see Material and methods). The lines a, b, c correspond to non stimulated cells, and the lines d, e, f correspond to cells stimulated with Zn SO₄.

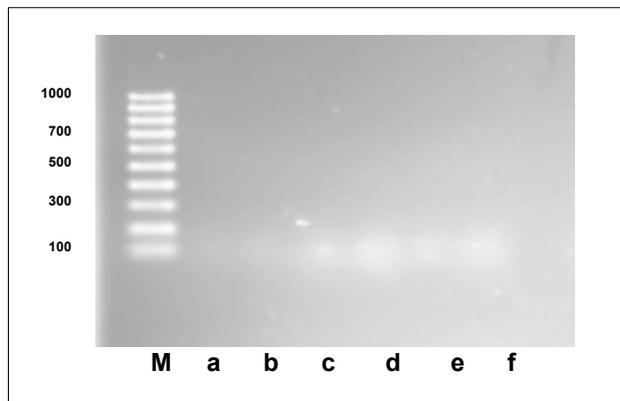


Figure 5b. C6 glioma cells transfected with pMT-Anti IGF-I. M- marker; a, b, c, d, e, f - amplified DNA using MHC-I primer "d" (see Material and methods). The lines a, b, c correspond to non stimulated cells, and the lines d, e, f correspond to cells stimulated with Zn SO₄. Note the expression of MHC-I demonstrated by RT PCR in cells corresponding to lines d, e, and f.

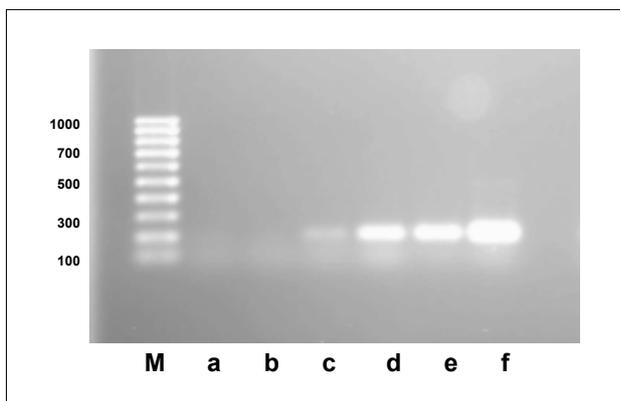


Figure 6a. C6 glioma cells transfected with pMT-Anti IGF-I-Receptor. Absence of expression of IGF-I demonstrated by RT PCR. M- marker; a, b, c, d, e, f - amplified DNA using IGF-I primer "b" (see Material and methods). The lines a, b, c correspond to non stimulated cells, and the lines d, e, f correspond to cells stimulated with Zn SO₄.

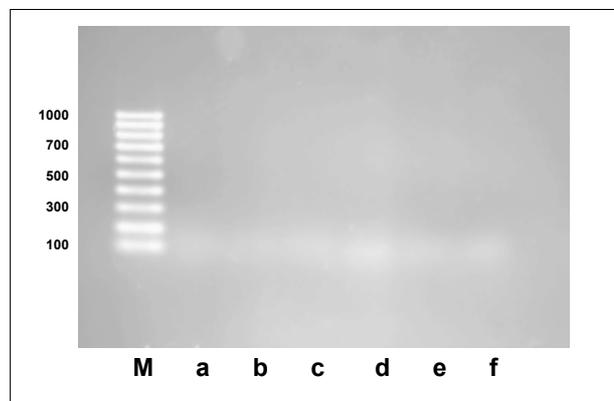
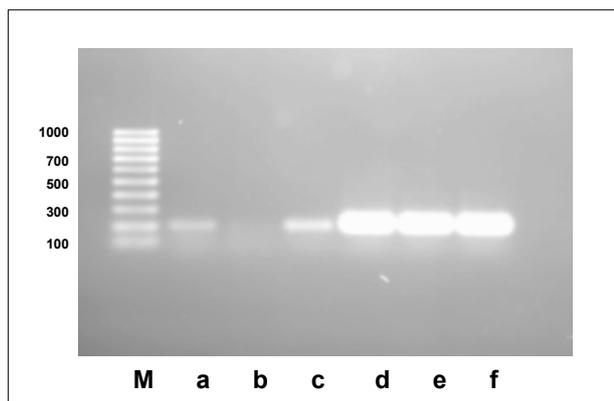


Figure 6b. C6 glioma cells transfected with pMT-Anti IGF-I-Receptor. M- marker; a, b, c, d, e, f - amplified DNA using MHC-I primer "d" (see Material and methods). The lines a, b, c correspond to non stimulated cells, and the lines d, e, f correspond to cells stimulated with Zn SO₄. Note the expression of MHC-I demonstrated by RT PCR especially in cells corresponding to lines d, e, and f and weakly in non stimulated cells (c).



gy for the neuroectodermal neoplasia - glioblastoma multiforme - have evolved, we have showed that a relationship may exist between the anti-tumor immune response and apoptosis. We demonstrated that IGF-I antisense transfection induce MHC-I molecules, and that using vectors encoding MHC-I antisense cDNA: 1) MHC-I expression, related to IGF-I antisense and triple helix transfection, can be reversible; 2) suppression of MHC-I expression in immunogenic IGF-I antisense transfected cells is following by the strike decrease of apoptotic cells [30]. These data define the immune and apoptotic characteristics of tumoral cells selected for cellular therapy of glioma, and permitted introduce the IGF-I antisense and triple-helix technology as a new gene therapy approach for treatment of human glioblastoma [23].

In here presented work down regulation of cellular IGF-I mRNA is accompanied by a decrease of cellular IGF-I. Transcription of the IGF-I gene is down regulated for 4 - 6 passages in rat glioma cells transfected with pMT-Anti IGF I-R vector. In

control experiments using the same vector carrying the lacZ gene in place of IGF-I antisense DNA sequence, we have showed 24 hours after transfection a 3-10% transgene frequency (100 % after three weeks) and strong cytoplasmic staining for X-gal in rat C-6 cells. The induction of IGF-I antisense RNA in transfected cells was followed by change in cell morphology, increase in apoptosis and enhanced expression of MHC-I. Transfected cells were long and narrow in shape and frequently string-like in appearance. This change was associated with down regulation in IGF-I protein. The change might be the signal either a reversion of the malignant phenotype or the recovery of some antigenic potential of these cells. As to the MHC-I expression, down-regulation of MHC-I due to action of IGF-I has been reported for experiments with rat thyroid cells [31]. This would be in agreement with results reported here concerning the inverse correlation between IGF-I and MHC-I protein expression in glioma cells. In tumor cells, the absence of IGF-I, when induced by IGF-I antisense technology, is associated with

massive apoptosis as previously demonstrated in transfected hepatoma cells [32]. In here presented work, the occurrence of morphological features vacuolization of cytoplasm suggests that the intracellular growth factor may be a control point for modulating to apoptotic process of glioma cells. IGF-I has been reported to block the apoptosis pathway in a variety of cell lines [33]. In contrast rats given injections of C6 glioma cells expressing antisense IGF-I-R did not develop tumors and were protected from a subsequent challenge with wild-type C6 glioma cells for at least months [3]. However, total eradication of pre-existing wild type tumors was not described [3]. On the other side, a qualitative relationship between the level of IGF-I receptor and tumorigenesis in nude mice, which correlates to the extent of apoptosis has been shown [34]. When the function of IGF-I receptor is decreased, glioma cells undergo massive apoptosis. It was concluded for the IGF-I-R result, that this receptor activated by its ligand plays a protective role against programmed cell death. This protection was even more striking in vivo than in vitro [34]. Another possible interpretation could be that an immune response occurring in the animals inhibits tumorigenesis. This is probably because nude mice do have a residual immune system containing both natural killer cells and B lymphocytes [35]. Progress in defining the pathophysiological role of signalling and downstream of IGF-I-R in neoplasia might lead to the development of novel targeting strategies [36]. Recent research has shown that targeting IGF-I-R leads to impressive antineoplastic activity in many in vitro and in vivo models of common human cancers. Strategies that have been used include [36]: administration of molecules to interfere with ligand binding to IGF-I-R, such as IGF-BPs, peptide or small-molecule competitive binding antagonists, or blocking anti-receptor antibodies [3, 37, 38], antisense or small interfering RNA strategies to reduce receptor expression [40]; introduction of a dominant-negative IGF-I-R to interfere with receptor action use of small-molecule IGF-I-R-specific tyrosine kinase inhibitors [41]; and targeting signalling pathways downstream of IGF-I-R with agents such as AKT or TOR inhibitors [42, 43]. The hypothesis that targeting the IGF-I-R will increase the efficacy of other antineoplastic treatments is based on evidence that survival signals originating at this receptor limit the efficacy of other treatments designed to induce apoptosis [36].

The results presented in this study show that inhibition of IGF-I-R up-regulates MHC-I expression in transfected "antisense" glioma cells. We need to add that increased expression of protease nexin I which may reduce the tumorigenic potential of the C-6 glioma cells was also observed when the IGF-I "triple-helix" cells or IGF-I receptor "triple-helix" cells were injected into nude mice [24, 27]. On the other hand, in our work concerning mouse hepatoma cells transfected by IGF-I triple-helix approach [28], we have observed the decrease of cytokines like IL-10, which is a strong immunosuppressor, and TNF- α , which can act as a factor stimulating a tumoral growth. Moreover we have found the increased level of TAP 1 and 2 in these cells. A further elucidation of the relationship between the immune process, related to MHC-I or HLA system [44], and the apoptotic process is under investigation. Our antisense IGF-I approach together with that of antisense IGF-I-R and other actually studied antisense strategies of gene therapies targeting bcl-2 [45], telomerase [46], urokinase-type plasminogen activator

receptor (uPAR) [47], neuroglia related Cell Adhesion Molecule [48] or matrix metalloproteinases [49] constitute the beginning of successful clinical trial to win "glioblastoma battle".

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