

# Treatment of glioblastoma multiforme cells with temozolomide-BioShuttle ligated by the inverse Diels-Alder ligation chemistry

Klaus Braun<sup>1</sup>  
 Manfred Wiessler<sup>1</sup>  
 Volker Ehemann<sup>2</sup>  
 Ruediger Pipkorn<sup>3</sup>  
 Herbert Spring<sup>4</sup>  
 Juergen Debus<sup>5</sup>  
 Bernd Didinger<sup>5</sup>  
 Mario Koch<sup>3</sup>  
 Gabriele Muller<sup>6</sup>  
 Waldemar Waldeck<sup>6</sup>

<sup>1</sup>German Cancer Research Center, Dept of Imaging and Radiooncology, Heidelberg, Germany; <sup>2</sup>University of Heidelberg, Institute of Pathology, Heidelberg, Germany; <sup>3</sup>German Cancer Research Center, Central Peptide Synthesis Unit, Heidelberg, Germany; <sup>4</sup>German Cancer Research Center, Dept of Structural Analysis of Gene Structure and Function, Heidelberg, Germany; <sup>5</sup>University of Heidelberg, Dept of Radiation Oncology, Heidelberg, Germany; <sup>6</sup>German Cancer Research Center, Division of Biophysics of Macromolecules, Heidelberg, Germany

**Abstract:** Recurrent glioblastoma multiforme (GBM), insensitive against most therapeutic interventions, has low response and survival rates. Temozolomide (TMZ) was approved for second-line therapy of recurrent anaplastic astrocytoma. However, TMZ therapy in GBM patients reveals properties such as reduced tolerability and inauspicious hemogram. The solution addressed here concerning GBM therapy consolidates and uses the potential of organic and peptide chemistry with molecular medicine. We enhanced the pharmacologic potency with simultaneous reduction of unwanted adverse reactions of the highly efficient chemotherapeutic TMZ. The TMZ connection to transporter molecules (TMZ-BioShuttle) was investigated, resulting in a much higher pharmacological effect in glioma cell lines and also with reduced dose rate. From this result we can conclude that a suitable chemistry could realize the ligation of pharmacologically active, but sensitive and highly unstable pharmaceutical ingredients without functional deprivation. The TMZ-BioShuttle dramatically enhanced the potential of TMZ for the treatment of brain tumors and is an attractive drug for combination chemotherapy.

**Keywords:** drug delivery, carrier molecules, facilitated transport, glioblastoma multiforme, temozolomide

## Introduction

The Grade IV classification of glioblastoma multiforme (GBM) means that these tumors grow aggressively and rapidly. The tumor entity GBM is considered one of the most aggressive and intractable cancers. Despite all treatments including neurosurgical resection followed by postoperative radiotherapy, the overall survival times of patients are unsatisfactory (Walker et al 1978). Therefore our aim is to use the potential for temozolomide (TMZ) to treat GBM by improving drug's potency. The combination of the oral administration of TMZ with radiotherapy led to longer survival times (Combs et al 2005) by delaying tumor progression. Including simple oral application increased the interest in using TMZ to treat GBM (Osoba et al 2000). The action of TMZ (8-carbamoyl-3-methylimidazo [5,1-d]-1,2,3,5-tetrazin-4(3H)-one) has been extensively studied, primarily in leukemia and lymphoma cells. TMZ is rapidly resorbed after oral application and spontaneously decomposes in aqueous solution at physiological pH to the cytotoxic methylating agent 5-(3-methyltriazene-1-yl)imidazole-4-carboxamide (MTIC) (Tsang et al 1990). The cytotoxicity of TMZ appears to be mediated mainly through adduction of methyl groups to O<sup>6</sup> positions of guanine (O<sup>6</sup>mG) in genomic DNA (Denny et al 1994) followed by recognition of this adduct by the mismatch repair system (MMR), which mispairs with thymine during the next cycle of DNA replication.

The half-life of TMZ (Riccardi et al 2003) in plasma and non-target-gene-specific alkylating mode of action can result in undesired adverse reactions which then hamper

Correspondence: Klaus Braun  
 German Cancer Research  
 Center (DKFZ), Dept of Imaging  
 and Radiooncology, Im Neuenheimer  
 Feld 280, D-69120 Heidelberg, Germany  
 Tel +49 6221 42 2495  
 Fax +49 6221 42 3375  
 Email k.braun@dkfz.de

the therapeutic outcome efficiency. For this reason new TMZ-derivatives were designed. The use of the peptide-based nuclear localization sequence (NLS) (Braun et al 2002; Heckl et al 2002), which leads to an active nuclear targeting, could minimize existing handicaps. Because of their higher molecular mass and their physico-chemical characteristics the transport across the cellular membrane of TMZ-NLS peptide conjugates is still poor. Therefore a carrier molecule is needed, which can – after the transmembrane passage into the cytoplasm – discharge its TMZ-NLS-Cys cargo so that a sufficient concentration of pharmacologically active molecules can reach their target site in the nucleus.

In this context numerous viral (Tabin et al 1982; Seymour 1992; Advani et al 2002; Conlon and Flotte 2004; Palmer and Ng 2005) and non-viral (Bangham and Papahadjopoulos 1966; Derossi et al 1996; Storm and Crommelin 1997; Vives et al 1997; Bourne et al 2000; Merdan et al 2002) well-documented transport vehicles were designed by different laboratories. They are able to facilitate the cellular transport of molecules with inappropriate physico-chemical properties for the transmembrane passage.

Our considerations for improving the transport of TMZ into the cell nucleus led to a search for suitable ligation modes of TMZ with a nuclear address peptide (a) which in turn is connected to carrier molecules (b) (Table 1).

A brief comment is required here for a better understanding of the following paragraphs. In chemistry the meaning of “ligation reaction” is the basis of Diels-Alder chemistry, on which we focus here, in contrast that in molecular biology, in which it means the ligase-catalyzed connection of nucleic acids at the 3'-hydroxy group and the 5'-phosphate terminus.

This first ligation (a) should permit a multi-modular connection of functional peptides and formation of higher peptide chain lengths compared to the different manufacturing techniques. It should also achieve a higher functional variability regarding the coupled peptides or substances in relation to mono-modular peptides generated by simple solid phase peptide synthesis or by recombinant chemistry. However the fast and selective covalent linkage of biologically active molecules under physiological conditions still represents a great challenge in the chemical praxis.

For this reason, such a ligation reaction should meet the following criteria:

(1) rapid course of the reaction, (2) independent of solvent properties, (3) no side reaction with other functional groups present in the molecule, (4) no special coupling-reagents, (5) an economical procedure, and (6) irreversible chemical

reaction characteristics. The following ligation methods are favorable:

### Enzymatic methods

The enzyme-mediated peptide ligation method is favorable to preserve native active structures of proteins and peptides (Anthony-Cahill and Magliery 2002). In this context the bacterial enzyme sortase (Mazmanian et al 1999) demonstrates an interesting approach: a transpeptidase that catalyzes a sequence-specific ligation of functional peptides harboring the ligand motif Gly-Gly-Gly to the amino terminus of the target molecule or peptide (Parthasarathy et al 2007).

### Chemical methods

The ‘Staudinger ligation’, which is based on the reaction of phosphines with azides under nitrogen elimination, is proposed as a further convenient candidate for ligation reactions (Kohn and Breinbauer 2004; Chandra et al 2006). The second chemical reaction useful for ligation is the thioester-method, which has been described by different groups (Camarero et al 2004; Yeo et al 2004). A method recently described demonstrates a ligation of  $\alpha$ -ketoacids and N-alkylhydroxylamines under production of CO<sub>2</sub> and water as by-products (Bode et al 2006). A further concept, the modular ‘click chemistry’, uses the most practical and reliable chemical transformations and could be considered a promising platform. Its applications are increasingly found in all aspects of drug discovery, ranging from offering valuable clues in combinatorial chemistry and target-templated in situ chemistry, to proteomics and DNA research, using bioconjugation reactions (Kolb et al 2001; Kolb and Sharpless 2003). In this concept the 1,3-dipolar cycloaddition, considered as ‘cream of the crop’ (Rostovtsev et al 2002), represents a prime example of the electrocyclic reactions developed by Huisgen (Huisgen 1984).

### Method used in our experiments

Our ligation approach is based on cycloaddition reactions via the pericyclic Diels-Alder reaction (DAR) with ‘inverse-electron-demand’, which is a modification of  $\pi$ -electron-deficient N-heteroaromatics with electron-rich dienophiles (Bachmann and Deno 1949). The main open question was how to circumvent the reversibility of conventional DAR. Here we used tetrazine-structures harboring electron-pure dienes permitting an irreversible reaction by formation of molecular nitrogen (Kim et al 2005).

We reveal how proper chemistry contributes to optimization of pharmacological properties of an already efficient drug such as TMZ, thereby possibly extending the spectrum of tumor entity treatments.

**Table 1** Schematic pattern of the TMZ-NLS and transport-peptides modules (modular schemata of the TMZ-BioShuttle)

Address-peptide with TCT <sup>*)</sup> -dienophile		
K(TCT)-NLS-C- module		
K(TCT)-KPKKRKVC <sup>**)</sup>	MW: 1250	<sup>*)</sup> $\text{I}_{\text{tetracyclo-[5.4.2.1.7.0.2.6.0.8.11]3,5-dioxo-4-aza-9,12-trideca-diene}}$ <sup>**)</sup> amino acids single letter code
Transmembrane transport-peptide		
CRQIKIWFQNRMMKKWKK <sup>**)</sup> MW: 2478		

The table depicts the transmembrane transport module (lower part of the table) and the nuclear localization sequence (NLS) connected with the modified dienophile component tetracyclo-[5.4.2.1.7.0.2.6.0.8.11]3,5-dioxo-4-aza-9,12-trideca-diene (TCT) via the  $\epsilon$ -amino coupling of the lysine (upper part of the table).

Because the bimodal radiochemotherapy with TMZ is classified as a standard treatment approach after primary diagnosis, we focused primarily on improving the established pharmaceutical TMZ by coupling it to the modular structured carrier, termed here as the TMZ-BioShuttle, and by testing its dramatically increased efficiency in different glioblastoma cell lines.

## Experimental

All reactions and procedures were carried out under normal atmospheric conditions. Table 1 shows the schematic

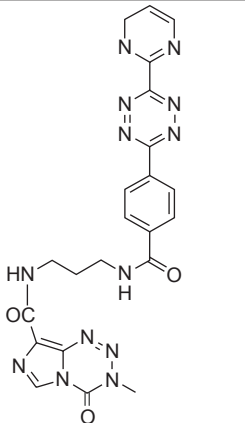
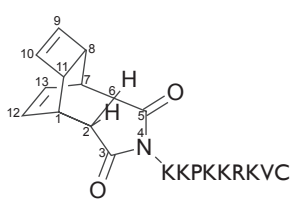
modules, Table 2 the educts, the intermediates and the products, and Table 3 the active substances investigated.

## Solid phase peptide synthesis of the K(TCT)-NLS-S-C-BioShuttle transporter

Syntheses of the transport module and the NLS module linked with lysine (TCT)

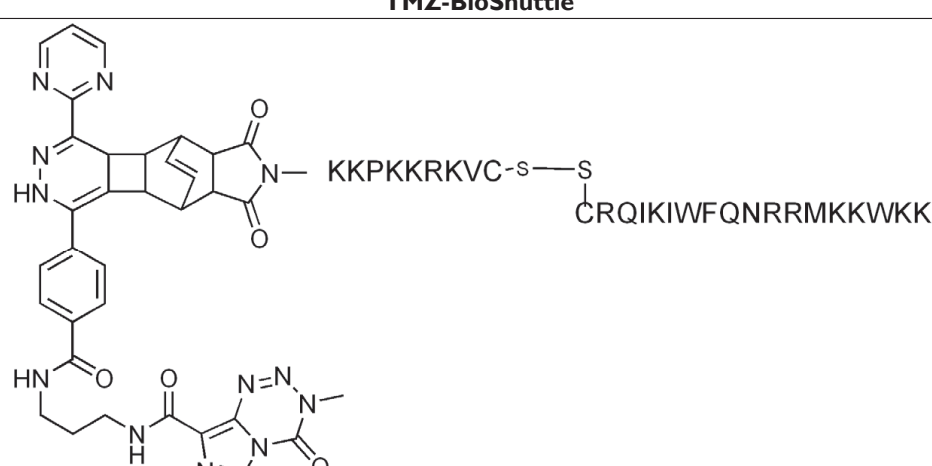
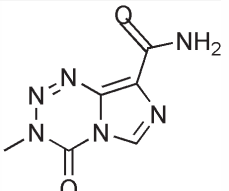
For solid phase synthesis of the K(TCT)-NLS-S-C-transport peptide we employed the Fmoc-strategy in a fully automated multiple synthesizer (Syro II) (Merrifield 1963). The synthesis

**Table 2** Educts for inverse-electron-demand-DAR and for the TMZ Cargo/K(TCT)-NLS-S-S-BioShuttle

TMZ-Cargo with Diene-Tetrazin-Spacer	Address-peptide with dienophile	Cleav. spacer	Transmembrane transport-peptide
		-S-S-	CRQIKIWFQNRRMKKWKK <sup>*)</sup>
MW: 513	MW: 3756		
<sup>*)</sup> Tetracyclo-[5.4.2 <sup>1,7</sup> .O <sup>2,6</sup> .O <sup>8,11</sup> ] <sup>**)</sup> 3,5-dioxo-4-aza -9,12-tridecadiene amino acids single letter code			

The left part of the table shows the amide-group modified temozolomide connected by epsilon-amino lysine to a tetrazine which acts as dien component.

**Table 3** Chemical structures of the investigated active alkylating substances

TMZ-BioShuttle

MW: 4242
Temozolomide (TMZ)

MW: 194

The chemical structure of the TMZ-BioShuttle is shown in the upper part of the table. The lower part shows the graphic formula of the alkylating agent temozolomide.

was carried out on a 0.05 mmol Fmoc-Lys(Boc)-polystyrene resin 1% crosslinked and on a 0.053 mmol Fmoc-Cys(Trt)-polystyrene resin (1% crosslinked). 2-(1H-Benzotriazole-1-yl)-1,1,3,3-tetramethyluronium hexafluorophosphate (HBTU) was used as coupling agent. The last amino acid of the NLS-peptide was incorporated as Boc-Lys(TCT)-OH. Cleavage and deprotection of the peptide resin were achieved by treatment with 90% trifluoroacetic acid, 5% ethanedithiol, 2.5% thioanisol, and 2.5% phenol (v/v/v/v) for 2.5 h at room temperature. The products were precipitated in ether. The crude material was purified by preparative HPLC on an Kromasil 300-5C18 reverse phase column (20 × 150 mm) using an eluent of 0.1% trifluoroacetic acid in water (A) and 60% acetonitrile in water (B). The peptides were eluted with a successive linear gradient of 25% B to 60% B in 40 min at a flow rate of 20 mL/min. The fractions corresponding to the purified protein were lyophilized.

#### Coupling of the transmembrane transporter- and the K(TCT)-NLS-C-modules

As a last step the K(TCT)-NLS-C and the transport peptide were oxidized in an aqueous solution of 2 mg/mL in 20% DMSO for about 5 hours. The oxidation progress was monitored by analytical C18 reversed-phase HPLC, and then peptide was purified as described above. The purified material

was characterized with analytical HPLC and laser desorption mass spectrometry in a Bruker Reflex II.

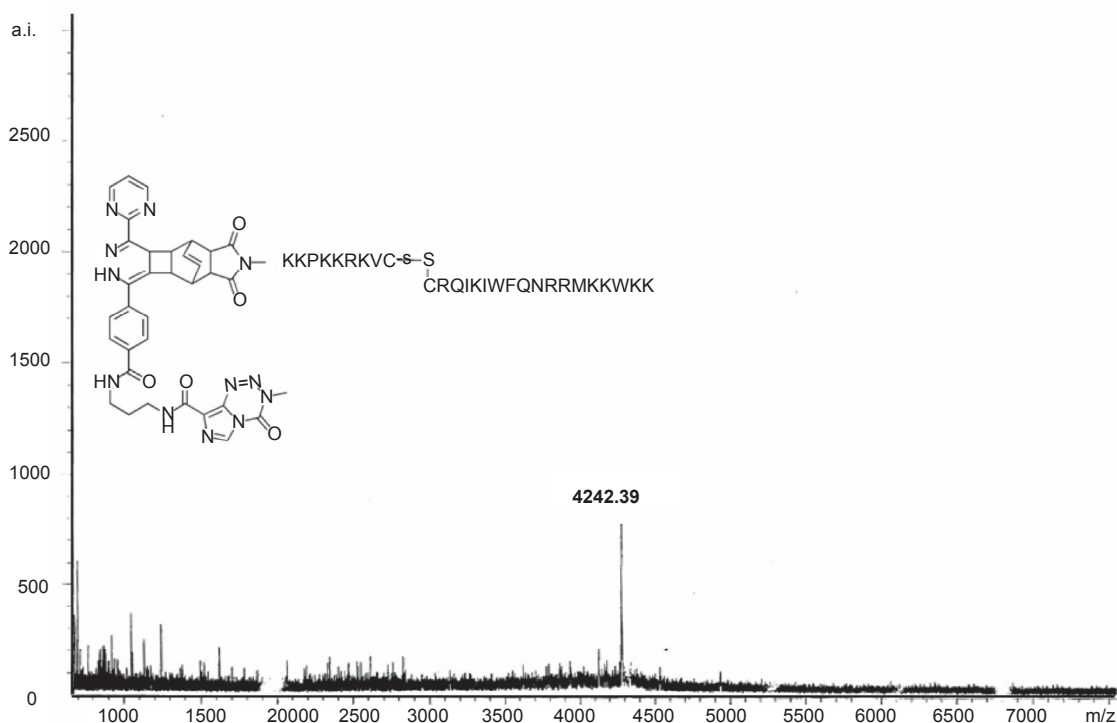
The empirical formula of the K(TCT)-NLS-C module is C<sub>64</sub>H<sub>110</sub>N<sub>18</sub>O<sub>12</sub>S with a MW 1355,74.

#### Ligation chemistry of the K(TCT)-NLS-S-S-BioShuttle and the TMZ-derivative

TMZ-<sub>(amino-bond)</sub>-tetrazolin-spacer linkage with the BioShuttle-carrier

A 0.2 mol preparation with 42.7 mg L1834 and 67.4 mg L1932 as well as 28 μL triethylamin were dissolved in chloroform. The reaction process runs undisturbed and provides a defined product with the molecular weight (MW) m/e 513. Further preparation: 0.3 mmol L1932 Triflat MW 445 and 0.4 mmol L1834 were dissolved in 100 μL triethylamin 0.7 mmol overnight at room temperature. Thin-layer chromatography (TLC) (9.5/0.5 chloroform/EtOH) indicates a defined product which points to the complete reaction.

TMZ-dien-spacer linked with K(TCT)-NLS-S-S-BioShuttle Preparation of 0.2 mmol (48 mg F77 and 44 mg L1834) with 0.2 mmol (0.03 mL) triethylamin: After dissolving the F77 in triethylamin and chloroform, the acid chloride of L1834 was added. The chemical reaction was carried out overnight at room temperature and the product was detected by TLC.



**Figure 1** Mass spectrometry plot of the TMZ-BioShuttle (as shown in Table 3) demonstrates the estimated molecular mass. The ordinate represents the absolute intensity (a.i.); the abscissa the mass to charge ratio (m/z).



By mass spectrometry we detected the mass of the chemical compound 4242.

## Reagents and cell culture

Human glioblastoma (GBM) primary cells (Gli98 and Gli366) were provided by the DKFZ division of Biophysics of Macromolecules. All cell lines were cultured in DMEM (Gibco Cat. No. 12800) supplemented with 10% FCS and maintained in culture at 37 °C with 5% CO<sub>2</sub> atmosphere and 95% humidity.

## Chemotherapy treatment

Pure temozolomide (TMZ) was purchased from Sigma-Aldrich, Germany (Cat. No. 76899) and the material was subdivided into two parts for subsequent processing. One part was followed up and coupled to the transporter molecules. As a control, the second part was dissolved in acetonitrile 10% (Sigma-Aldrich, Germany) with a final concentration of 0.2% acetonitrile. Further control studies with acetonitrile were performed to exclude potential toxic effects of acetonitrile.

Gli98 and Gli366 cells were suspended ( $1 \times 10^5$  cells/mL) in DMEM (control) and in DMEM containing appropriate amounts of TMZ and the TMZ-BioShuttle (50 μM equimolar) and cell behavior was investigated for up to 6 days of induction time.

## Cell cycle and cell death analysis

The effects on cell viability and cell cycle distribution were determined by DNA flow cytometry. Flow cytometric analyses were performed using a PAS II flow cytometer (Partec, Muenster, Germany) equipped with a mercury vapor lamp (100 W) and a filter combination for 2,4-diamidino-2-phenylindole (DAPI) stained single cells. From native probes the cells were isolated with 2.1% citric acid/0.5% Tween 20 according to the method for high resolution DNA and cell cycle analyses (Ehemann et al 2003) at room temperature with slight shaking. Phosphate buffer (7.2 g Na<sub>2</sub>HPO<sub>4</sub> × 2H<sub>2</sub>O in 100 mL H<sub>2</sub>O dest.) pH 8.0 containing DAPI was used for staining the cell suspension. Each histogram represents 30,000 cells for measuring DNA-index and cell cycle. For histogram analysis, we used the Multicycle program (Phoenix Flow Systems, San Diego, CA).

### Cell viability – assessed by flow cytometry with the PI-method

For detection of apoptotic cells and viability, a FACS Calibur flow cytometer (Becton Dickinson Cytometry Systems, San Jose, CA) was used with filter combinations for

propidium iodide. The Cellquest program (Becton Dickinson Cytometry Systems, San Jose, CA) was used for analyses and calculations. Each histogram and dot plot represents 10,000 cells. After preparation according to Nicoletti et al (1991) with modifications (Singer et al 2007; Tschaharganeh et al 2007), measurements were acquired in the logarithmic mode in FI-3 and calculated by setting gates over the first three decades to detect apoptotic cells. Dead cells are positive for propidium iodide and stained red; living cells remain unstained. In the logarithmic histogram the positions of unstained living cells are in the first 2 decades, the 3rd decade contains cells with membrane damage, and dead cells are placed in the 4th decade.

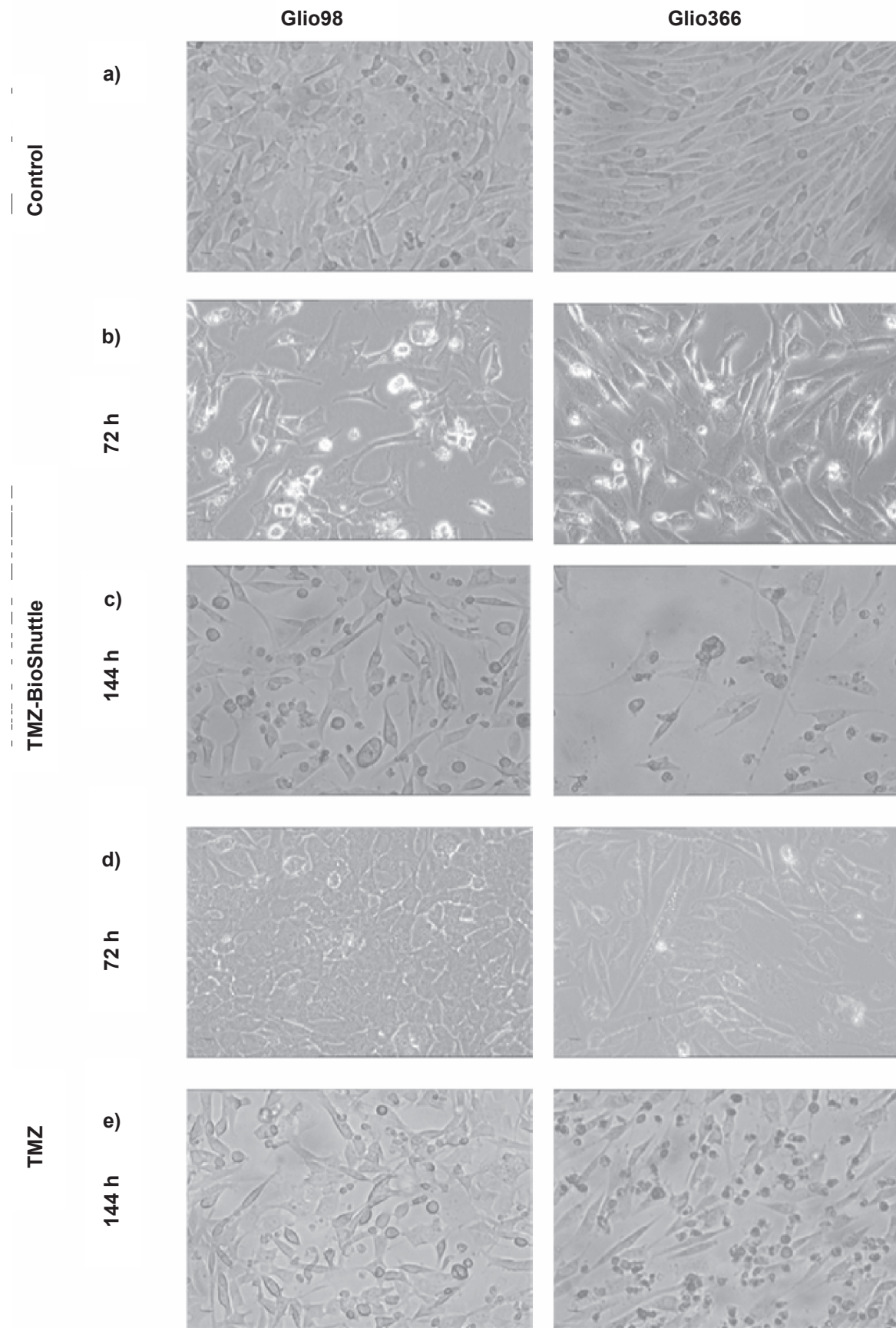
## Results

### Time-course of cell growth and cell death in Gli98 and Gli366 cells induced by TMZ and TMZ-BioShuttle – pharmacological effects

Two different glioma cell lines, herein referred as Gli98 (Figure 2a-e, left column) and Gli366 (Figure 2a-e, right column) were used to investigate the effect of TMZ and TMZ-BioShuttle on brain tumor cells. After 24 hours of treatment with the TMZ-BioShuttle, no change in the phenotype of both glioma cell lines could be observed under the light microscope (data not shown); however in the cell culture medium an increasing number of aggregated cells could be observed. In our opinion after 72 hours of treatment with TMZ cells a possible reconstitution of the phenotype was shown and also a regrowth 144 hours after TMZ treatment (Figure 2e). The cells treated with the TMZ-BioShuttle seemed not to regenerate.

After 144 hours of treatment with the TMZ-Bioshuttle an increase of aggregated Gli366 cells and of smaller scale aggregations in Gli98 cells in the cell culture medium were observed (not visible in the picture). Both adherent cells types reveal a non-confluent state because of many dead cells in the TMZ-treated probes. The cell density declines dramatically from 72 to 144 hours after TMZ-BioShuttle treatment in both glioma lines but the decrease of adherent cells was greater in Gli366 cells than in Gli98 cells (Figure 2c).

To learn more about the cell death in the two treated glioma cells after treatment, we carried out comet assays in parallel probes (compared to Figure 2). Basically the Gli98 and Gli366 cells did not exhibit fragmented DNA; no comets of the Gli98 cells 72 hours after treatment with TMZ (Figure 3d) and TMZ-BioShuttle (Figure 3b) can be



Figures 2a–e Microscopical monitoring of the treated glioblastoma cells.

observed and they resembled the untreated Glioblastoma control cells (Figures 3a in both cell lines).

After 144 hours of treatment with TMZ (Figure 3e) the glioblastoma cells show different-sized and fragmented dead cells and comet structures. The shape of the two observed comets was not used for calculation since it was not a major event. The TMZ-BioShuttle treated cells are mainly swollen (Figure 3c). Furthermore, the total cell density of the TMZ-only treated samples was lower, compared with that in control and TMZ-BioShuttle treated samples. The comet assay study detected no DNA fragmentation in the Glioblastoma98 and the Glioblastoma366 until 72 hours after treatment. 144 hours after application of TMZ-BioShuttle large amounts of cell fragments could be observed in all treated cell lines (Figure 3c). DNA fragments could be detected in Glioblastoma366 (Figure 3e right column) but comets in the Glioblastoma98 cells are not visible 144 hours after TMZ treatment (Figure 3e left column).

### Cell cycle study

As can be seen in the next experiment the predominant event was discovered in the cell cycle investigation. Flow cytometry is considered to be a valuable tool for high throughput measurements of the cell cycle, cellular dynamics, cell activities and cell toxicity studies. It provides sensitive and specific data on the cellular behavior of glioblastoma cells after treatment with TMZ and the TMZ-BioShuttle conjugate. We determined the cell cycle behavior of glioma cells, for example, of the Glioblastoma366 cells after drug treatment. To investigate the influence of TMZ with and without BioShuttle transporter we performed a time-course analysis of the cell cycle. Cell growth after drug treatment was dramatically different. The pharmacological effect on brain tumor cells was shown 144 hours after treatment with TMZ-BioShuttle.

Figure 4 shows cell cycle distribution of Glioblastoma366 cells 144 hours after treatment: The plots named control, TMZ-BioShuttle, and TMZ are shown from left to right. The cell cycle distribution is indicated as G1-phase, S-phase and the G2/M phase. These cells show a diploid cycle (red colored). The plot of the untreated control is shown (Figure 4, left) and exhibits a G1 cell fraction of 79% and a G2 cell fraction of 10%. The contingent of cells in the S-phase amounts to 11%. Compared to the S-phase fraction of untreated control the percentage of S-phase cells treated with TMZ is decreased to 8%. The TMZ-BioShuttle treated Glioblastoma366 cells clearly exhibit a decreased amount of cells in the S-phase (2%). Compared to control, the TMZ and the TMZ-BioShuttle treated probes display an increased cell number in the G2/M phase of 20% and 72%, respectively. The cells in the G1-phase of the

TMZ probe decreased slightly (72%) whereas in cells in the G1-phase of the TMZ-BioShuttle treated Glioblastoma366 cells decreased significantly (26%) as shown in Table 4. The amount of debris (blue colored) is visible and each histogram represents 30,000 measured cells.

### Cell viability study on human Glioblastoma98 Glioblastoma366 cells

The cell killing effect of TMZ and the TMZ-BioShuttle on the glioblastoma cell lines Glioblastoma98 and Glioblastoma366 is demonstrated in Figure 5 and in the corresponding values implemented in the Table 5.

The pharmacological effect of the TMZ-BioShuttle on Glioblastoma366 and Glioblastoma98 cells differs greatly from that of the non-derivatized TMZ.

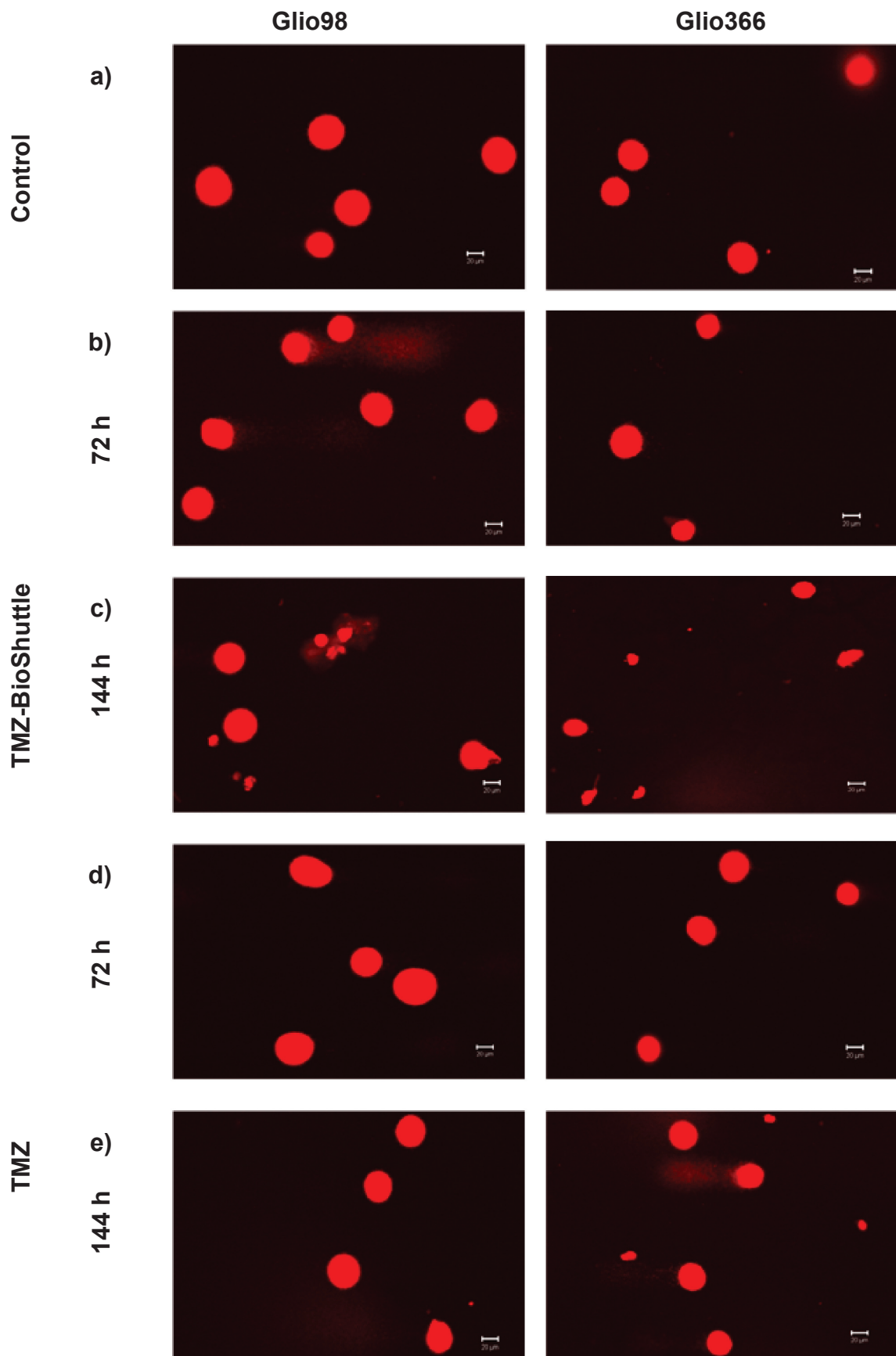
The cell lines show a different sensitivity to chemotherapy with TMZ and TMZ-BioShuttle respectively. After 72 hours of treatment with TMZ-BioShuttle, 28% of Glioblastoma366 cells were dead; after 144 hours of TMZ-BioShuttle treatment 32.5% of the Glioblastoma366 cells were dead. The effect of TMZ on the Glioblastoma366 cell line was less noticeable compared to control. Dead cells at 72 hours after TMZ treatment are in the range of untreated controls (Glioblastoma366: 4.5% increases to 7%) after 6 days. The percentage of dead cells after TMZ treatment was decreased slightly (by 3.5%). 144 hours after TMZ treatment no difference was detectable between percentage of dead control cells and treated cells (7.5%). The percentage of dead cells is an underestimation because many cells were already degraded (Figure 2c).

The sensitivity of the investigated Glioblastoma98 cells to TMZ and TMZ-BioShuttle is somewhat different from that of Glioblastoma366 cells. Percentage of dead Glioblastoma98 control cells increased from 2.5% (3 days) to 10.5% (6 days). Three days after TMZ treatment, the relative amount of dead cells barely differed from the corresponding control cells (3.5%). Six days after TMZ treatment the amount of dead cells increased to 14.5%. After 3 days of TMZ-BioShuttle treatment Glioblastoma98 cells exhibit 13.5% dead cells, and 22% after 6 days of treatment.

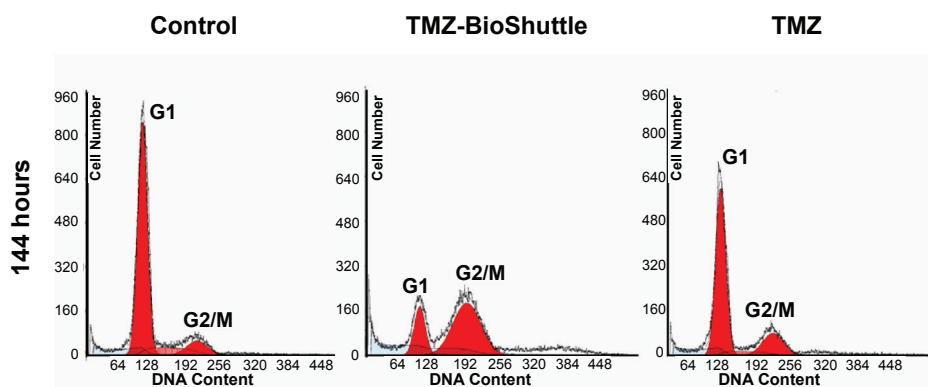
## Discussion

Treatment of high-grade gliomas poses a substantial challenge to neuro-oncologists. The main problems are positions of the tumors in the cerebrum, which allow a radical excision only in a limited number of patients, and a high recurrence rate after excision. Current available chemotherapy (nitrosoureas, procarbazine and vincristine or the PCV protocol) is now at least 25 years old and even when combined with radiotherapy has not changed the fatal prognosis for patients





**Figure 3a–e** Comet assay studies of untreated cells, treated with TMZ and TMZ-BioShuttle for different cell lines; the scale bar represents 20  $\mu$ m.



The figure shows FACS scans of Glioblastoma cells treated for 144 hours with TMZ-BioShuttle (middle) and TMZ (right). The left part of the picture displays the cell cycle distribution of the untreated control cells.

Debris cells      Cells in the G1 and G2/M phases

**Figure 4** Cell cycle of Glioblastoma cells after application of TMZ and TMZ-BioShuttle.

The figure shows FACS scans of Glioblastoma cells treated for 144 hours with TMZ-BioShuttle (middle) and TMZ (right). The left part of the picture displays the cell cycle distribution of the untreated control cells. Debris cells      Cells in the G1 and G2/M phases.

with high-grade glioma; also, progression-free survival in patients with glioblastoma multiforme has been greatly improved. Therefore, the initial reports of a substantial antitumor activity of temozolomide (TMZ) in brain tumors demonstrated by Newlands et al could indicate a relief (Newlands et al 1992). TMZ, a DNA-methylating agent of the imidazotetrazine class, was developed in the 1980s, through rational drug design (Denny et al 1994; Newlands et al 1997).

With regard to undesired side-effects like the often-observed strong suppression of the peripheral lymphocytes, the connection of the ‘old fashioned’ TMZ drug as a cargo to transporter molecules (TMZ-BioShuttle) was investigated. These peptides, designed for facilitating rapid transport across cellular membranes, could improve their delivery and their targeting under the pharmacological activity in cells and tissues protecting cells in the bloodstream.

The treatment of glioblastoma Glioblastoma cells with TMZ interferes with the cell cycle and results in a decreased number of cells in the S-phase fraction as shown 144 hours after TMZ application (11% in the control and 8% in the TMZ treated cells). Furthermore, in the TMZ-BioShuttle treated cells the smallest contingent of the S-phase cells of 2% was detectable (Figure 4 and Table 4) after 144 hours.

The fact that the glioblastoma cells arrest in the G2/M-phase after TMZ-treatment was documented by Hirose et al (2003) and could not be confirmed by the tested Glioblastoma cells, but the treatment with TMZ-BioShuttle clearly increased cell number residing in the G2/M-phase by up to 72%. The increase seems to be attributable to interactions

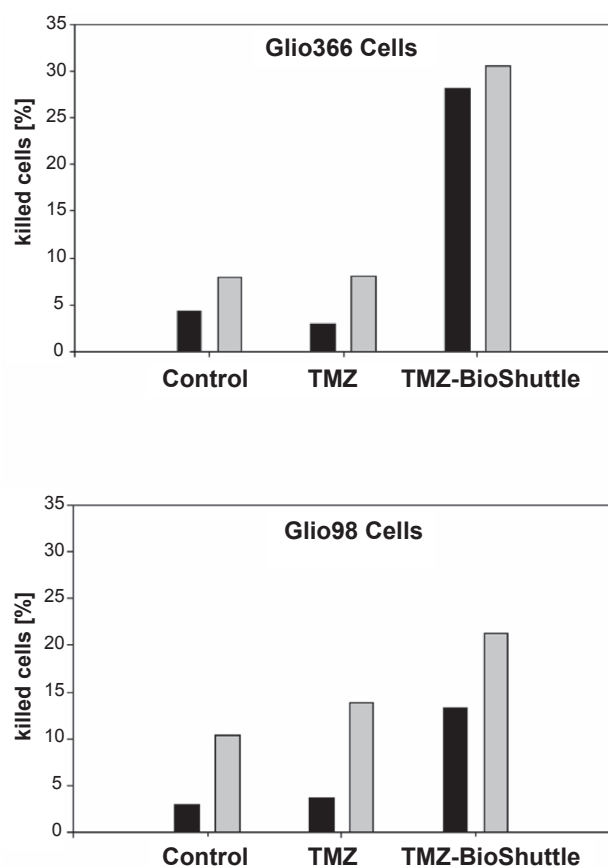
of TMZ and TMZ-BioShuttle with the mitogen-activated protein (MAP) kinase p38 $\alpha$ , which is activated by the mismatch repair system (MMR) and is responsible for the TMZ-induced G2/M block (Hirose et al 2003).

The phenomenon of a slightly reduced S-phase (8% versus control 11%) in the TMZ treated Glioblastoma cells is not contradictory to the common effects of alkylating agents which retard rate of cell division (Bignold 2006). A possible explanation for the strongly decreased S-phase cell number in the TMZ-BioShuttle treated probes would be the existence of a S-phase cell cycle arrest as documented in a bimodal TMZ/interferon- $\beta$  (IFN- $\beta$ ) study (Park et al 2006). Our flow cytometry experiments also showed a strong cell cycle arrest in the S-phase (Thomas et al 1996). A further criterion for a prolonged late S/G2-phase suggests an implication of the histone acetyltransferase 1 (HAT1) which participates in recovering block-mediated DNA damage (Barman et al 2006).

**Table 4** Relative amount of cells in the cell cycle 72 and 144 hours after TMZ and TMZ-BioShuttle treatment

	Cell fraction	Control	TMZ-BioShuttle	TMZ
72 hours	G1 [%]	73	33	74
	G2 [%]	19	17	16
	S [%]	8	50	10
144 hours	G1 [%]	79	26	72
	G2 [%]	10	72	20
	S [%]	11	2	8

The table is derived from the FACS scan data and shows an overview of the cell cycle distribution of Figure 5 calculated as a percentage.



**Figure 5** Flow cytometric estimation of cell killing of Gliob366 (upper part) and of Gliob98 cells (lower part). The abscissae of the graph show the relative amount of the dead Gliob366 cells after 72 hours (black bars) and after 144 hours (gray bars) after exposure to the TMZ-BioShuttle (right bar-pair). The bars at the left site indicate the untreated control cells and the bar pair in the middle, the dead cells, TMZ-treated. The ordinates represent the relative amount of dead cells (%).

After exposure to the TMZ-BioShuttle for 144 hours, the observed strong increase of the G2-phase cells of Gliob366 cells could possibly originate from the documented inhibition of the RNA and protein syntheses, which are necessary for the successful completion of G2 and the initiation of mitosis. These G2-arrested cells were found to be deficient in certain proteins that may be specific for the G2-mitotic transition (Rao 1980). Obstruction of the cell cycle process results in the decrease of the G1-phase cell fraction (26%) of the TMZ-BioShuttle treated Gliob366 cells, which appears

to be a consequence of their disability passing in the mitotic process.

The fact that TMZ is also known to induce a transient proliferation arrest combined with the accumulation of cells in the G2/M phase could be confirmed as shown in the decrease of the cells in the S-phase and in turn by the increase of cells in the G2/M-phase from 10% (control) to 20%. In parallel a decrease of the G1-phase cells (72%) compared to the control (79%) could be observed also in the Gliob366 after 6 days. The conditioning to the cell cycle of the untreated control cells and cells 6 days after TMZ application could be a process described as potentially dependent on the p53 status (Hirose et al 2001). It is documented that in glioma, p53 wild-type cells undergo prolonged cell cycle arrest associated with elevation of p53 levels, and although the cells are non-proliferative, they remain viable. Conversely, p53-deficient cells undergo a more transient cell cycle arrest and within 12 days lose viability by mitotic catastrophe and/or apoptosis. Preclinical models demonstrated a schedule-dependent antitumor activity against high-grade glioma, suggesting increased activity with repeated exposure to TMZ (Raymond et al 1997).

Further, the activity of the O<sup>6</sup>-methylguanine-DNA-methyltransferase (MGMT) is involved in the DNA-repair-procedures in cells. The epigenetic silencing of the MGMT DNA-repair gene is associated with longer survival in glioblastoma patients and benefits from alkylating agents like TMZ (Hegi et al 2005). Possibly after treatment, the cell cycle behavior of the Gliob366 could be explained with epigenetic properties like CpG island hypermethylation of the DNA repair enzyme MGMT. The restricted TMZ activity and the crucial role of the repair enzyme MGMT are documented in the literature (Antoun et al 2000; Walton et al 2008). Here we did not consider to which extent the enrichment of pharmacological DNA-active methylating substances like TMZ could contribute to overcoming the opposite effects. If and to what extent MGMT effects are involved remains to be seen in further studies.

Figure 5 shows a dramatically increased sensitivity of both the Gliob98 and the Gliob366 cells to TMZ compared

**Table 5** Relative amount of dead Gliob366 and Gliob98 cells [%]

	After 72 hours			After 144 hours		
	control	TMZ	TMZ-BioShuttle	control	TMZ	TMZ-BioShuttle
Gliob366	4.5	3.5	28	7.5	7.5	32.5
Gliob98	2.5	3.5	13.5	10.5	14.5	22

Actual amount of killed cells derived from Figure 5.

with the TMZ-BioShuttle. The present data document a different cellular contumaciousness to TMZ treatment. A real effectiveness of the pure TMZ in Glio366 and Glio98 cells could not be observed, whereas clear cell killing effects caused by TMZ-BioShuttle were detected in contrast.

It is important to note that the TMZ-BioShuttle treatment of Glio366 and Glio98 glioma cells redounds to cell killing effects as shown in Figure 5 and Table 5. In the TMZ treated Glio366 probes dead cells barely differed from their ratio in the untreated cells after 3 days (3%–4%) as well as after 6 days (9%) after TMZ application, but the percentage of dead cells treated with TMZ-BioShuttle escalated to 28% after 3 days and 32.5% after 6 days.

Furthermore the novel properties of the TMZ-BioShuttle could give reason to the enlargement of other tumor entities like the therapy of prostate tumors.

The application of the TMZ-BioShuttle could minimize the handicap of TMZ for the therapy of patients with brain tumors but the establishment of the TMZ-BioShuttle necessitates new ways for the synthesis. Conditions that hampered the above postulated criteria such as rapid and whole concurrent chemical reactions in aqueous solution at room temperature for a proper chemical ligation of functional peptides could be circumvented using the ‘inverse-electron-demand’ of the Diels-Alder reaction. This reaction enhances the economics of the chemical reaction by the following parameters: the increase in the reaction rate, gentle reaction conditions at room temperature and the resulting reaction kinetics demand no excess at all in the educts. Furthermore this careful execution of the chemical reaction could realize the ligation of pharmacologically active but sensitive and highly unstable pharmaceutical ingredients without functional deprivation. The potential of the TMZ modified by coupling to the BioShuttle carrier (TMZ-BioShuttle) has been confirmed for the treatment of brain tumors and remains an attractive drug for further clinical combination chemotherapy.

## Dedication

This article is dedicated to Harald zur Hausen, the winner of the Nobel Price Award 2008 in Medical Sciences, and the retired Head of the German Cancer Research Center.

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## Abbreviations

DAR, Diels-Alder-reverse; GBM, glioblastoma multiforme; MGMT, methylguanine-DNA-methyltransferase; NLS, nuclear localization sequence; TCT, tetracyclo-[5.4.2<sup>1,7</sup>.O<sup>2,6</sup>.O<sup>8,11</sup>]3,5-dioxo-4-aza-9,12-tridecadien; TMZ, temozolomide.

## Disclosures

The authors declare no conflicts of interest.

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