

1 **On optimal temozolomide scheduling for slowly growing gliomas**

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51 **Background:** Temozolomide (TMZ) is an oral alkylating agent active against
52 gliomas with a favorable toxicity profile. It is part of the standard of care in the
53 management of glioblastoma, and is commonly used in low-grade gliomas. *In-*
54 *silico* mathematical models can potentially be used to personalize treatments and
55 to accelerate the discovery of optimal drug delivery schemes.

56

57 **Methods:** Agent-based mathematical models fed with either mouse or patient
58 data were developed for the *in-silico* studies. The experimental test beds used to
59 confirm the results were: mouse glioma models obtained by retroviral expression
60 of EGFR wt or EGFR vIII in primary progenitors from p16/p19 ko mice and grown
61 *in vitro* and *in vivo* in orthotopic allografts, and human glioblastoma U251 cells
62 immobilized in alginate microfibers. The patient data used to parametrize the
63 model were obtained from the TCGA/TCIA databases and the TOG clinical study.

64

65 **Results:** Slow growth 'virtual' murine gliomas benefited from increasing TMZ
66 dose separation *in silico*. In line with the simulation results, improved survival,
67 reduced toxicity, lower expression of resistance factors and reduction of the tumor
68 mesenchymal component were observed in experimental models subject to long-
69 cycle treatment, particularly in slowly-growing tumors. Tissue analysis after long-
70 cycle TMZ treatments revealed epigenetically-driven changes in tumor
71 phenotype, which could explain the reduction in glioma growth speed. *In-silico*
72 trials provided support for methods of implementation in human patients.

73

74 **Conclusions:** *In-silico* simulations, and *in-vitro* and *in-vivo* studies show that
75 TMZ administration schedules with increased time between doses may reduce

76 toxicity, delay the appearance of resistances and lead to survival benefits

77 mediated by changes in the tumor phenotype in gliomas.

78

79 **Keywords:** Glioblastoma, Mathematical Oncology, Temozolomide, Drug

80 resistance, Persisters, *In-silico* trials, optimal drug scheduling, personalized

81 oncology, tumor phenotype.

82

83 **IMPORTANCE OF THE STUDY**

84

85 *In-vivo* evidence is provided of improvements in survival, resistance, and toxicity
86 from TMZ schemes with long rest periods between doses in slowly-growing GBM
87 mouse models. The results match hypotheses generated *in silico* using a
88 mathematical model incorporating the main biological features and fed with real
89 patient data. An epigenetically-driven change in tumor phenotype was also
90 revealed experimentally, which could explain the reduction in glioma growth
91 speed under the 'long cycle' scheme. To determine the extent to which our results
92 hold for human patients, large sets of simulations were performed on virtual
93 patients. These *in-silico* trials suggest different ways to bring the benefits
94 observed in experimental models into clinical practice.

95

96 INTRODUCTION

97 Adult gliomas are the most common primary malignant tumors of the central
98 nervous system. A novel classification of these neoplasms has been proposed¹⁻
99 ³, using a combination of molecular and histopathological features,. Especially
100 important for patient prognosis is the presence of mutations in the isocitrate
101 dehydrogenase 1/2 (IDH1/2) genes, which distinguishes IDH mutant (IDHmut)
102 gliomas from IDH wild-type (IDHwt) glioblastomas (GBM)⁴. GBMs are diagnosed
103 at a later age (median 64) and are characterized by the presence of additional
104 alterations. Some of them are mutations in the telomerase reverse transcriptase
105 (TERT) promoter, amplification of the epidermal growth factor receptor (EGFR),
106 and gains and losses of chromosomes 7 and 10, respectively⁵. GBMs have a
107 dismal prognosis (15 months' overall survival) despite the standard-of-care
108 treatment, which consists of maximal surgical resection followed by radiotherapy
109 (RT) plus concomitant and adjuvant chemotherapy (CT) with temozolomide
110 (TMZ), an oral alkylating agent^{6,7}. TMZ is administered orally at a dose of 75
111 mg/m² daily throughout RT, plus six cycles of maintenance TMZ 150–200 mg/m²
112 for 5 out of 28 days⁶. The cytotoxicity of this drug is attributed to the addition of
113 methyl groups to DNA, and especially to the formation of O6-methylguanine (O6-
114 meG) lesions and the subsequent formation of double-strand breaks during DNA
115 replication, which requires cell division for the emergence of the cytotoxicity. As
116 O6-meG can be removed by methylguanine methyltransferase (MGMT) in tumors
117 expressing this protein, MGMT promoter methylation is considered a predictive
118 biomarker of TMZ response in gliomas⁸. However, challenges remain in
119 establishing reliable inter-laboratory assays⁹, as well as in estimating the effect
120 of limited MGMT promoter methylation on outcomes¹⁰. Moreover, based on the

121 short half-life of TMZ, it was suggested that high doses or repeated doses could
122 improve the effect of this commonly-used drug, and reduce the capacity of cells
123 to repair the DNA¹¹. However, dose-dense TMZ, with the administration of lower
124 but continuous doses with the aim of depleting intracellular MGMT, did not show
125 improved efficacy in newly diagnosed GBM,¹² and has shown only modest results
126 in recurrent tumors¹³⁻¹⁶ at the cost of increased hematological toxicity. Simple
127 mathematical models of lower grade gliomas (LGG) (grade 2/3 gliomas) have
128 suggested that TMZ schemes with longer rest periods between cycles could
129 improve the survival of patients¹⁷. However, no study has tested the efficacy of
130 longer spacing between cycles/doses. Therefore, there may be room for
131 improvements in the schedules used for TMZ administration to glioma patients.
132 Mathematical models describe real systems by abstraction and mathematical
133 formalism. They enable extrapolation beyond the situations originally analyzed,
134 allowing for quantitative predictions, inference of mechanisms, falsification of
135 underlying biological hypotheses, and quantitative descriptions of relationships
136 between different components of a system. They cannot replace experimental
137 results obtained by biomedical models, but may complement experimentation in
138 providing a broader picture¹⁸. Moreover, in the field of oncology they can suggest
139 what the best RT, CT or combination regimens might be, aiding the
140 implementation of the treatment of different cancers, including gliomas. A number
141 of mathematical models of LGGs have been constructed in order to study the
142 optimal delivery scheme of cytotoxic therapies^{17,19,20,21}. It is important to
143 emphasize that in slow-growth gliomas, such as LGGs, only a small percentage
144 of cells is proliferating as shown by fractions of Ki67 positive cells typically below
145 5%²². Notably, even in GBM, there is a small group of tumors with reduced

146 Ki67%²³. It may be guessed that intensive therapies intended to deliver the
147 maximum tolerated dose in the shortest possible time may be overkill for these
148 tumors. Indeed, several authors have proposed that schemes with longer spacing
149 between doses could produce better results in LGG patients^{17,20,21,24}. However,
150 the simple mathematical models developed previously account only for a limited
151 number of key biological processes. Notably, no detailed theoretical models,
152 including realistic resistance mechanisms, have been considered previously.
153 Also, it is still unknown whether the potential gain observed *in silico* for LGGs
154 would apply to GBMs as well. First of all, the proneural (PN) to mesenchymal
155 (MES) phenotypic transition observed in GBMs, either spontaneously or as a
156 result of (CT/RT), would limit the effect of TMZ as the tumor becomes more
157 resistant^{25,26}. Also, from the biological point of view, there is a growing body of
158 literature suggesting that the evolution of tumor cells to a fully drug-resistant state
159 may often proceeds through a reversible drug-tolerant phase²⁷, so-called
160 persister cells²⁸. Rabé et al identified a population with persister characteristics
161 under TMZ treatment of glioma cells²⁹. One may guess that longer spacing
162 between cycles could delay the PN-MES transition, as well as allowing for
163 persister cells to revert to their normal sensitive states. Neither of these facts has
164 been accounted for by previous mathematical modeling approaches.

165 The intention of this paper was to set out a proof of concept supporting treatment
166 regimes with longer times between doses of TMZ (hereafter referred to as
167 protracted temozolomide schedules, PTS). Our hypothesis was that PTS could
168 lead to reduced appearance of persister cells. Moreover, reduced toxicity and
169 increased tolerance was anticipated from increasing the rest period between
170 cycles. To test these ideas, a stochastic mesoscopic discrete mathematical

171 model was first produced, including all relevant biological processes expected to
172 play a role in the response of GBMs to TMZ. Next, PTS was studied *in vitro* and
173 in animal models and found to be in good agreement with the *in silico*
174 observations. The tissue of TMZ-responding allografts was then analyzed, and
175 showing a striking change in tumor phenotype, with an increase in PN and a
176 decrease in MES markers, driven by epigenetic changes, which could explain the
177 reduction in tumor growth. Finally, exploratory virtual 'clinical' trials were
178 performed with *in-silico* tumors simulated with the previous mathematical model,
179 to guide the implementation of the concept in clinical practice. The results
180 indicated that survival is increased as spacing between doses becomes
181 progressively larger, until a threshold is reached; spacings beyond this threshold
182 would fail to improve survival.

183

184 **METHODS**

185

186 **Cell lines and cell culture**

187 The human GBM U251 cell line was purchased from American Type Culture
188 Collection (ATCC, USA). It was cultured in Dulbecco's Modified Eagle Medium
189 (Biological Industries, USA) supplemented with 10% fetal bovine serum (Sigma-
190 Aldrich, Germany), 2 mM glutamine (Sigma-Aldrich, Germany), 5,000 U/ml
191 penicillin, and 5 mg/ml streptomycin (Gibco™, Thermo Fisher Scientific, United
192 States). Cells were cultivated at 37°C in a humidified 5% CO₂ atmosphere and
193 passaged twice a week after reaching 80-90% confluence using 0.25%
194 trypsin/EDTA.

195

196 Mouse SVZ cell lines were obtained by retroviral expression of EGFRwt or
197 EGFRvIII (pBabe-EGFR wt (#11011) and MSCV-XZ066-GFP-EGFR vIII
198 (#20737)) in primary neural stem cell cultures obtained from the subventricular
199 zone (SVZ) of p16/p19 ko mice³⁰. After infection, the cells were injected into nude
200 mice, and the tumors that grew were dissociated and the lines SVZ-EGFRwt/amp
201 and SVZ-EGFRvIII were established³¹. Both models express GFP and luciferase
202 as a reporter. Cells were grown as previously described^{30,31}. Briefly, they were
203 maintained in stem cell medium; Neurobasal (Invitrogen) supplemented with B27
204 (1:50) (Invitrogen); GlutaMAX (1:100) (Invitrogen); penicillin-streptomycin (1:100)
205 (Lonza); 0.4% heparin (Sigma-Aldrich); and 40 ng/ml EGF and 20 ng/ml bFGF2
206 (Peprotech). For dissociation and passage Accumax (ThermoFisher) was used.

207

208 **Production of alginate microfibers with U251 immobilized cells**

209 Alginate microfibers with cells were produced by extrusion as described earlier³².
210 U251 cell lines were immobilized in alginate microfibers by the same procedure.
211 Briefly, 4×10^6 cells/ml were mixed with a 2 % w/v Na-alginate solution to obtain
212 final concentrations of 1.5% w/v Na-alginate. The Na-alginate solution with cells
213 was manually extruded through a blunt edge stainless steel 25G needle
214 immersed in the gelling bath (3 % w/v $\text{Ca}(\text{NO}_3)_2 \times 4\text{H}_2\text{O}$). Due to the exchange of
215 Na^+ with Ca^{2+} , the liquid stream solidified in the gelling bath, thus forming
216 insoluble microfibers. The microfibers were left in the bath for 15 min in order to
217 complete gelling and were then washed with medium. After cell immobilization,
218 0.5 g of alginate fibers were distributed into a T25 flask and cultured for 28 days
219 without passage in 13 ml of MEM medium. 50% of the medium was changed
220 twice a week.

221

222 **Viability study**

223 The impact of three different TMZ (Sigma-Aldrich) (100 μ M) treatment modalities
224 (everyday treatments, X+1, and protracted (every 3 days, X+3; every 7 days,
225 X+7)), starting from day 7, was determined by comparing the effects on cell
226 viability, morphology, and aggregation using a CalceinAM (CAM)/propidium-
227 iodide (PI) assay. U251 cells immobilized in alginate microfibers were cultured
228 for 28 days and stained using CAM/PI as a LIVE/DEAD staining. Alginate
229 microfibers containing cells were incubated for 45 min at 37 °C in medium with
230 CAM in a final concentration of 4 μ M while PI was added to a final concentration
231 of 5 μ M. Fluorescence microscopy images were taken using a Leica TCS SP5 II
232 Basic confocal laser-scanning microscope (Leica Microsystems CMS GmbH;
233 Germany), visualizing live (green) and dead (red) cells at every z-axis
234 encompassing the alginate microfiber. Z-stack projections and quantitative
235 estimation of the cell mass were analyzed using ImageJ software.

236

237 ***In-vitro* treatments of mouse cells**

238 SVZ EGFR wt/amp cells were incubated in the presence of TMZ (25 μ M), which
239 was supplemented three times: one day (X+1), three days (X+3) or seven days
240 (x+7) after the first dose. In a different experiment, SVZ EGFR wt/amp cells were
241 treated with TMZ (25 μ M) and/or Azacytidine (AZA) (Sigma-Aldrich) (5 μ M) for 8
242 days. For both experiments, cells were collected and lysed and subsequently
243 analyzed by qRT-PCR, as described below.

244

245 **Intracranial tumor formation and treatment *in vivo***

246 Animal experiments were reviewed and approved by the Research Ethics and
247 Animal Welfare Committee at “Instituto de Salud Carlos III” (PROEX 02/16), in
248 agreement with the European Union and national directives. Intracranial
249 transplantation to establish orthotopic allografts was performed injecting 300,000
250 cells (resuspended in 2 μ l of culture cell medium) with a Hamilton syringe into
251 athymic Nude-Foxn1nu brains (Harlan Iberica). Female mice (2-3 months of age)
252 were used, 7 to 10 animals per group. The injections were made into the striatum
253 (coordinates: A–P, –0.5 mm; M–L, +2 mm, D–V, –3 mm; related to Bregma) using
254 a Stoelting Stereotaxic device. Mice were treated with TMZ (10 or 50 mg/kg
255 through intraperitoneal injection, i.p., with the schedules given in the various
256 experiments). TMZ was dissolved in PBS+1% BSA, which was used to treat
257 control animals.

258

259 **Immunohistochemical (IHC) staining**

260 Tumor samples were fixed in 10% formalin overnight, dehydrated through a
261 series of graded ethanol baths and then infiltrated with paraffin. Then, 2.5 μ m-
262 thick sections were obtained in a microtome and then sections were rehydrated
263 and permeabilized (1% triton X-100). Antigen retrieval was performed with Citrate
264 Buffer (10 mM, pH 6) in a pressure cooker (2 min). Endogenous peroxidase
265 inhibition and blocking with normal horse serum was also carried out before the
266 incubation with primary antibodies (anti-rabbit caspase3, 1:100, Cell signaling
267 #9662), anti-mouse ki67, 1:100 Dako #M7248) (overnight, 4 °C) and biotinylated
268 secondary antibodies (HRP anti-mouse and HRP anti-rabbit, 1:200, GE
269 Healthcare (2h at room temperature). Target proteins were detected with the ABC
270 Kit and the DAB kit (Vector Laboratories).

271

272 **Western Blot analysis**

273 For protein expression analysis, mouse tumor tissue was processed by
274 mechanical disruption in a lysis buffer (Tris-HCl pH 7.6, 1mMEDTA, 1mMEGTA,
275 1% SDS, and 1% Triton X-100) followed by heating for 15 min at 100°C. Protein
276 content was quantified by using a BCA Protein Assay Kit (Thermo Fisher
277 Scientific). Approximately 30 µg of proteins were resolved by 10% or 12% SDS-
278 PAGE, and these were then transferred to a nitrocellulose membrane (Hybond-
279 ECL, Amersham Biosciences, Little Chalfont, UK). The membranes were blocked
280 for 1 h at room temperature in TBS-T (10 mM Tris-HCl (pH 7.5), 100 mM NaCl,
281 and 0.1% Tween-20) with 5% skimmed milk, and then incubated overnight at 4
282 °C, with the corresponding primary antibody (mouse anti-MGMT 1:1000, BD
283 Biosciences, #557045), mouse anti-GAPDH (1:1.500, Santa Cruz Biotechnology
284 #sc-47724), rabbit anti-pTyr1068-EGFR (1:1.000, Cell Signaling #3777) and
285 rabbit anti-phospho-NF-kB p65 (Ser536) (1:1000, Cell Signaling #3033) diluted
286 in TBS-T. After being washed 3 times with TBS-T, the membranes were
287 incubated for 2 h at room temperature with their corresponding secondary
288 antibody (HRP-conjugated anti-mouse (#NA931) or anti-rabbit (#NA934),
289 Amersham Biosciences) diluted in TBS-T.

290

291 **RNA extraction and RT-PCR**

292 The impact of three different TMZ treatment modalities was determined by
293 comparing the effects on gene expression in human glioma cells and in the
294 mouse models.

295 - Human glioma cells: After 28 days of incubation, alginate microfibers with
296 U251 cells were dissolved and cells were released. To release cells, alginate
297 microfibers were dissolved in 0.5 mM EDTA for 10 min at 37°C. Total RNA
298 was extracted from control and treated group of cells. The extractions were
299 carried out using Tri Reagent Solution (Invitrogen LifeTechnologies, USA)
300 according to the manufacturer's instructions. cDNA was synthesized using 2
301 µg total RNA and High-capacity cDNA reverse transcription kit (Applied
302 Biosystems, USA) according to the manufacturer's instructions.

303 - Mouse samples. Brain tumors were dissected out after the mouse sacrifice
304 and fresh frozen. Alternatively, mouse cells grown *in vitro* were collected and
305 fresh frozen. RNA was extracted from the tissue or the cells using the RNA
306 isolation Kit (Roche). Total RNA (1µg) was reverse transcribed with
307 PrimeScript RT Reagent Kit (Takara).

308 Quantitative real time PCR was performed using the Light Cycler 1.5 (Roche)
309 with the SYBR Premix Ex Taq (Takara). The primers used for each reaction are
310 indicated in Supplementary Table S1. All experiments were performed in triplicate
311 and relative gene expression levels were analyzed by the 2^{-ddCt} method³³.

312

313 ***In-silico* analysis**

314 The Cancer Genome Atlas (TCGA) GBM dataset was accessed via UCSC xena-
315 browser (<https://xenabrowser.net>) to extract proliferation gene expression levels.

316 Classification into classical, mesenchymal, neural and proneural subtypes was
317 retrieved from the TCGA GBM data set⁵, together with gene expression values.

318 Differences in gene expression between different groups were calculated using
319 Student's t-test.

320

321 **Statistical analysis**

322 The difference between experimental groups was assessed by paired *t*-test and
323 one-way analysis of variance (ANOVA). For Kaplan-Meier survival curves, the
324 significance was determined by the two-tailed log-rank test. All analyses were
325 carried out with the GraphPad Prism 5 software. *P* values below 0.05 were
326 considered significant (**P* < 0.05; ***P* < 0.01; ****P* < 0.001; *****P* < 0.0001; n.s.,
327 not significant), both for mouse and simulated tumors. All experimental
328 quantitative data presented are the means +/- SEM from at least three samples
329 or experiments per data point.

330

331 **Discrete mathematical model**

332 An on-lattice agent-based mesoscopic model³⁵ was adapted to simulate the
333 longitudinal growth dynamics of glioma and its response to treatments *in silico*. A
334 comprehensive model description is provided in the Supplementary Information.
335 This study included three basic cellular populations: proneural cells (either
336 proliferative PNs or quiescent PNq), persister cells (P), and mesenchymal cells
337 (either proliferative MESs or quiescent, MESq). The cell dynamics between
338 different compartments is summarized in Figure S4 (and in the Supplementary
339 Information). It was assumed that PN cells may become MES cells, either directly,
340 due to local vessel damage and hypoxia once the local cell density exceeds a
341 critical threshold, or through a transient intermediate persister state induced by
342 TMZ exposure. Both of these routes were associated with the emergence of TMZ
343 resistance, as MES cells are assumed to be more resistant to TMZ than PN cells.

344 The methodology was first used to run simulations to explore the influence of the
 345 parameters on outcome, and to test the efficacy of the use of different dose
 346 spacings using murine parameters. Murine tumors were simulated without
 347 treatment (control), and treated with 3 TMZ doses, separated by 1, 4, 7 and 13
 348 days. Human tumors were simulated without treatment, under standard TMZ
 349 therapy (6 cycles of TMZ given for 5 days, and resting periods of 3 weeks), and
 350 under two different courses of TMZ, increasing the spacing between doses. A
 351 first set of simulations was performed by increasing the rest periods between
 352 doses from 3 weeks to 9 weeks. Another set of computational studies was run by
 353 increasing the spacing between individual doses, while removing the rest periods.
 354 In particular, spacings of 8 and 12 days between doses were considered. Human
 355 and murine parameters were estimated from previous studies (see Table 1) and
 356 our own datasets. Virtual human simulations were fed with real patient data^{35,36}
 357 to generate realistic tumors *in silico*.
 358 The simulator was implemented in Julia (version 1.1.1). Simulation file processing
 359 and graphics were undertaken in MATLAB (R2021a, MathWorks). Simulations
 360 were performed on two 2.4 GHz, 16-core, 192 GB memory Mac Pro machines.
 361 Computational cost per murine simulation ranged from 5 to 10 minutes, while for
 362 humans, the computational cost ranged from 20 to 50 minutes per simulation.
 363

PARAMETER	MEANING	VALUE (MOUSE)	VALUE (HUMAN)	UNITS	REFERENCE
τ_P	Division time (proliferative cells)	19	48-72	h	Estimated from [29]
ρ_M	Migration coefficient	0.12	0.4	mm ² /day	Estimated from [29]

τ_D	Death time	100	288-312	h	Estimated from [29]
μ_{SQ}	Transition rate from proliferative state to quiescent	0.3333-1	0.2-0.6	day ⁻¹	Estimated from [29]
μ_{QS}	Transition rate from quiescent state to proliferative	0.0166-0.1	0.03-0.2	day ⁻¹	Estimated from [29]
μ_{SP}	Transition rate from PNs to persister	0.0333	0.0333	min ⁻¹	Estimated from [27]
μ_{PS}	Transition rate from persister to PNs	0.2	0.2	day ⁻¹	Explored in this work
μ_{PR}	Transition rate from persister to MESs	0.0111	0.0111	min ⁻¹	Estimated from [27]
μ_{PT}	Transition rate from PNs to MESs	0.0208-0.0833	0.0208	h ⁻¹	Explored in this work
S_F	Survival fraction after TMZ dosing	0.75	0.5	unitless	Estimated from [38]

364

365 **Table 1:** Model parameters used to run murine and human simulations.

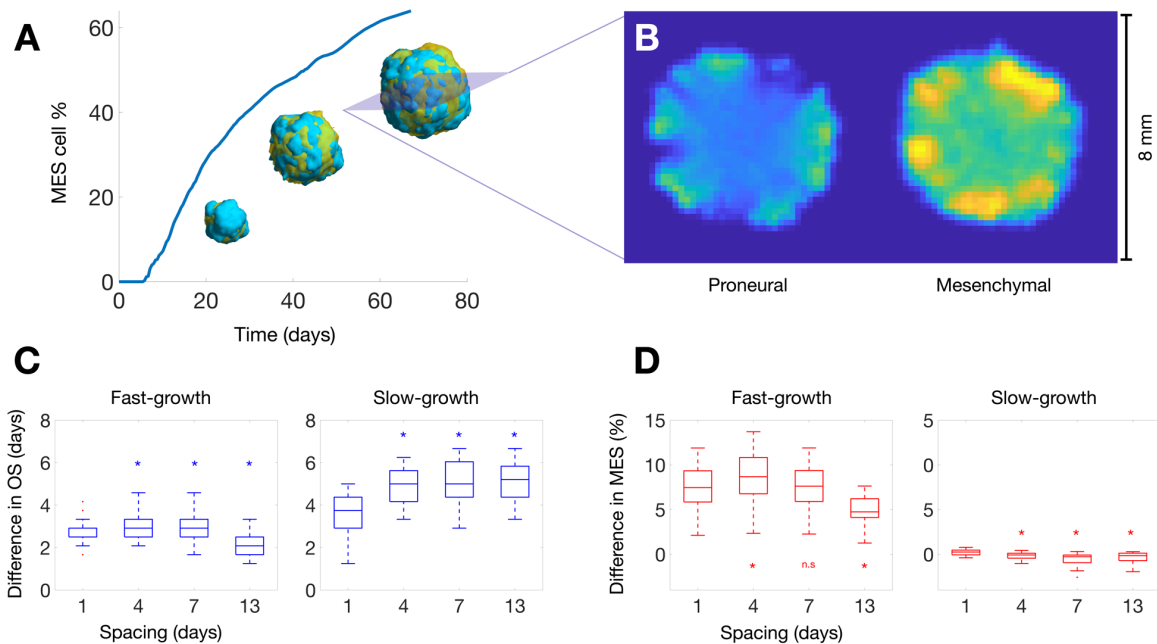
366

367 RESULTS

368 **Slow growth murine gliomas benefitted from increasing TMZ dose**
 369 **separation *in silico*.**

370 Virtual murine tumors were simulated as described in Methods, to explore the
 371 effect of different TMZ schemes in OS and MES content. A sustained increase in
 372 MES cell abundance was observed in untreated tumors (Figure 1A), reproducing
 373 the aforementioned PN-MES transition. These values are in agreement with real

374 mouse data obtained in *in-vivo* experiments performed in this study. Notably, the
375 tumor boundary was specially enriched in MES cells (Figure 1B).
376 Treatment with TMZ yielded an increase in OS for both fast- and slow-growth
377 tumors (Figure 1C). This survival increase was higher for slow-growth tumors;
378 moreover, enlarging the spacing between doses produced a better response in
379 this kind of tumors. Regarding resistance, TMZ induced a significant increase in
380 MES cell content in fast-growth tumors (Figure 1D). However, slow-growth
381 tumors did not undergo such increase; most of them remained with the same
382 amount of MES cells as control tumors, or even reduced their MES levels. This
383 effect was more evident for long spacings.
384 A robust observation was that virtual mice with slow-growth murine tumors had
385 the largest survival increase when increasing the spacing between doses *in silico*.
386 This benefit was preserved through the regions of the parameter space explored.
387 Longer spacing between doses led to reduced mesenchymal component in the
388 final tumors as compared to the 1-day spacing. Altogether, the simulations
389 suggest that longer spacing between doses would be more effective against slow-
390 growth gliomas, both in terms of OS and resistance development (Figures 1C-
391 1D).



392

393 **Fig. 1.** Mice tumor simulations predict an improvement in antitumor effect of TMZ
 394 in slow-growth GBMs. **(A)** Depiction of PN-MES transition in a single simulation,
 395 showing the evolution of MES cell abundance over time. 3D volumes are
 396 rendered at 50, 75 and 100% of simulation time (blue: PN cells; orange: MES
 397 cells). **(B)** Axial plane showing cell number per voxel in a simulated tumor at the
 398 end of simulation. Both PN and MES cell numbers belong to the same slice. **(C)**
 399 OS gain (in days) produced by TMZ dose spacing compared against control. 4-,
 400 7- and 13-day spacings were compared against 1 day spacing (Wilcoxon signed-
 401 rank test). **(D)** MES cell increase (percentage) compared against control. 4-, 7-
 402 and 13-day spacings were compared against 1 day spacing (Wilcoxon signed-
 403 rank test, *P <= 0.05, n.s.=non-significant).

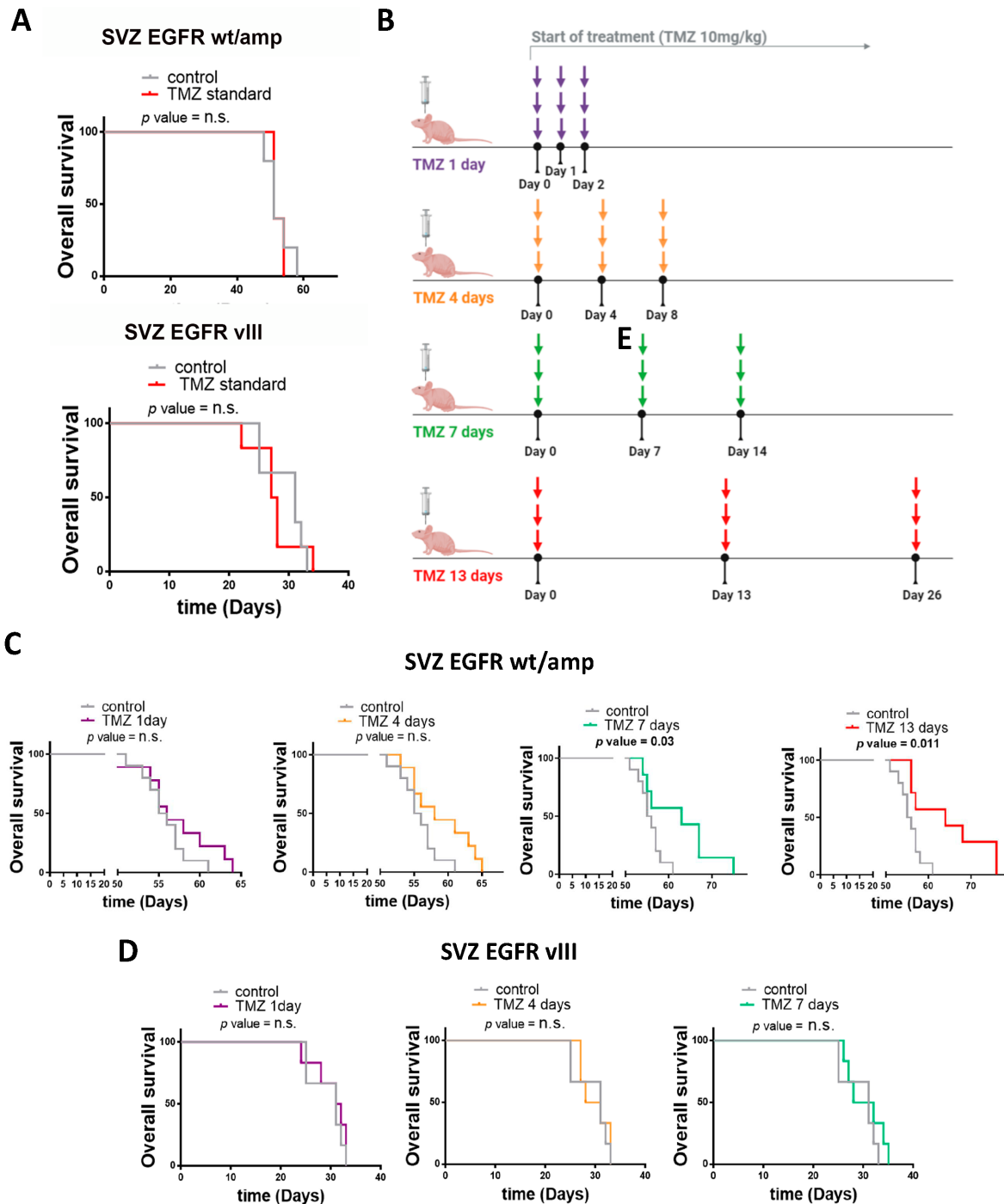
404

405 **Effect of protracted TMZ in two mouse glioma models with different**
 406 **proliferation kinetics.**

407 To validate the hypothesis that protracted TMZ regimes could offer therapeutic
 408 advantages depending on the degree of proliferation of glioma cells, mouse
 409 models were used, generated in our lab by overexpressing EGFRwt or EGFRvIII
 410 in p16/p19 ko subventricular zone (SVZ) progenitors. Both models generate
 411 gliomas in nude mice with a high penetrance and reproducibility. Notably, animals

412 survive for two months after SVZ-EGFRwt cell intracranial injection, whereas
413 SVZ-EGFRvIII tumors kill the animals much faster³¹, in agreement with the higher
414 aggressiveness attributed to the mutated isoforms of EGFR^{40,41}. Moreover, our
415 previous analyses showed that tumors formed by SVZ-EGFRvIII cells were much
416 more proliferative than those formed by SVZ-EGFRwt cells³¹.

417 Standard treatment of mouse glioma models with TMZ consisted of daily (5
418 days/week) i.p. injections of 10 mg/kg/day of the compound, which did not
419 produce a survival benefit in animals bearing SVZ-EGFRwt or SVZ-EGFRvIII
420 tumors (Figure 2A). Several TMZ regimes were then designed with three
421 consecutive doses of three TMZ injections, separated by 1 day (X+1), 4 days
422 (X+4), 7 days (X+7) or 13 days (X+13) (Figure 2B), although this last schedule
423 could not be applied in the EGFRvIII model due to their faster growth. The graphs
424 in Figure 2C show a clear reduction in SVZ-EGFRwt tumor growth in the X+7 and
425 X+13 protracted schemes. Extending the interval between TMZ doses did not
426 improve the response of SVZ-EGFRvIII bearing animals (Figure 2D). This result
427 shows the truth (at least in mice) of the suggestion from the mathematical model
428 that slowly-growing tumors are those that most benefit from protracted regimes.



429

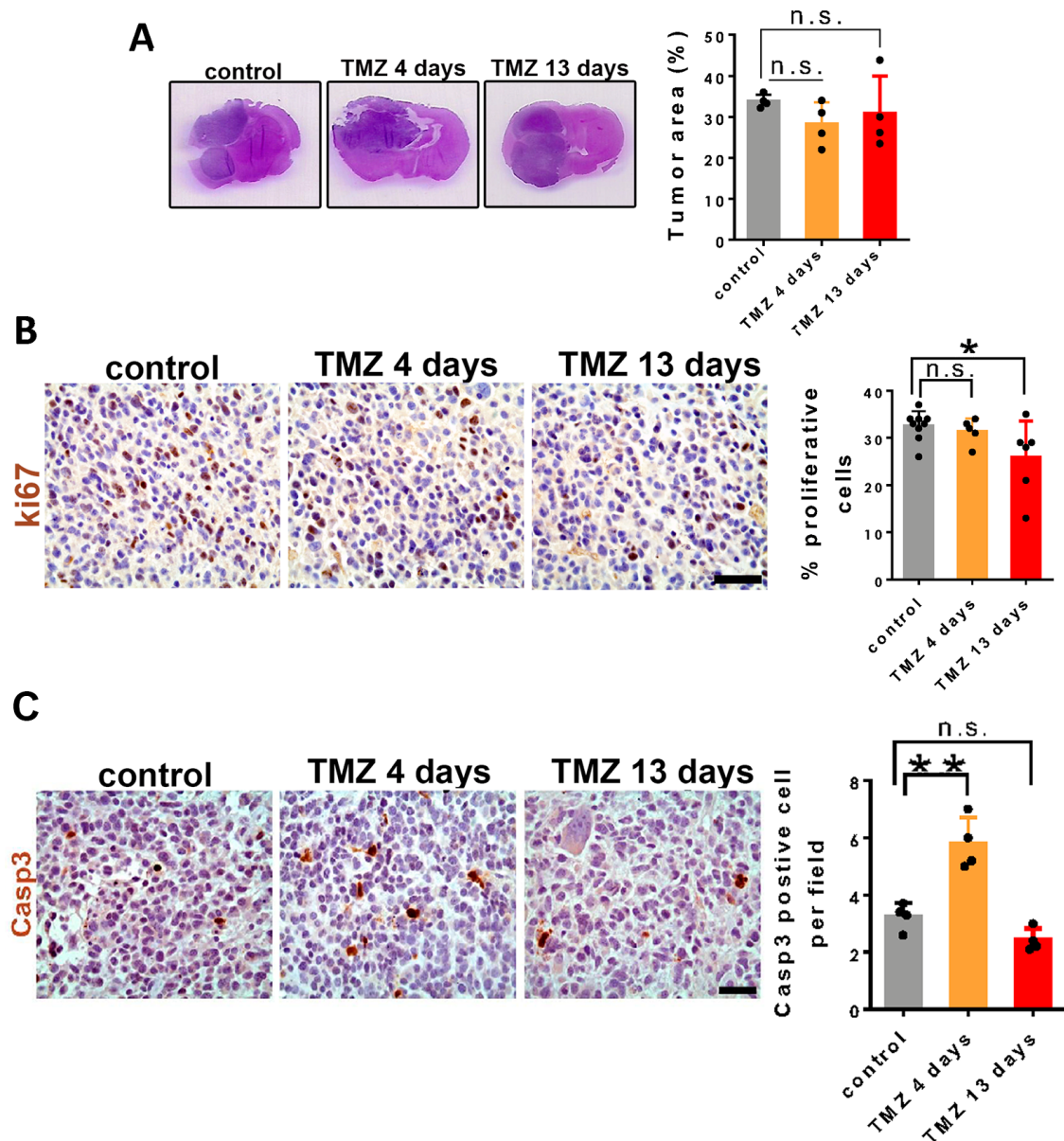
430 **Fig. 2.** Increasing spacing between TMZ doses improves the anti-tumor effect of
 431 TMZ in the SVZ-EGFR wt/amp (lower growth rate) but not in the SVZ EGFR vIII
 432 (faster growth speed) model. (A) Kaplan-Meier overall survival curves of mice that
 433 were orthotopically injected with SVZ EGFR wt/amp (top) or SVZ EGFR vIII
 434 (bottom) cells and subsequently treated with intraperitoneal injections (five days
 435 per week) of temozolomide (TMZ) (10 mg/kg per day) ($n = 6$). (B) Representative
 436 scheme of the different TMZ schedules studied. (C-D) Kaplan-Meier overall

437 survival curves of mice that were orthotopically injected with SVZ EGFR wt/amp
438 (n = 9) (C) or SVZ EGFR vIII (n = 6) (D) cells and subsequently treated with
439 intraperitoneal injections of TMZ (10 mg/kg per day) following the different
440 treatment protocols explained in (B), each of them represented by its assigned
441 color.

442

443 **Increasing spacing between TMZ doses does not increase cell death in the**
444 **tumors but reduces the expression of persister-related genes.**

445 To understand the beneficial effect of PTS, tumor tissue in the SVZ-EGFRwt
446 model was analyzed, comparing the control-treated with the X+4 (no response)
447 and the X+13 (responsive) tumors. Notably, no changes were found in the tumor
448 size in any of the schemes (Supplementary Figure 1A). In the X+13 scheme, a
449 small decrease was observed in the number of proliferating cells (Supplementary
450 Figure 1B), with no increase in the number of apoptotic cells (Figure 3A)
451 compared to control tumors. However, in the X+4 scheme there was no change
452 in proliferation, whereas an increase in the number of apoptotic cells was
453 measured (Supplementary Figure 1C). These results suggest that the reduction
454 in the tumor growth observed after X+13 protracted administration of TMZ is not
455 mediated by an increase in the cell death response.



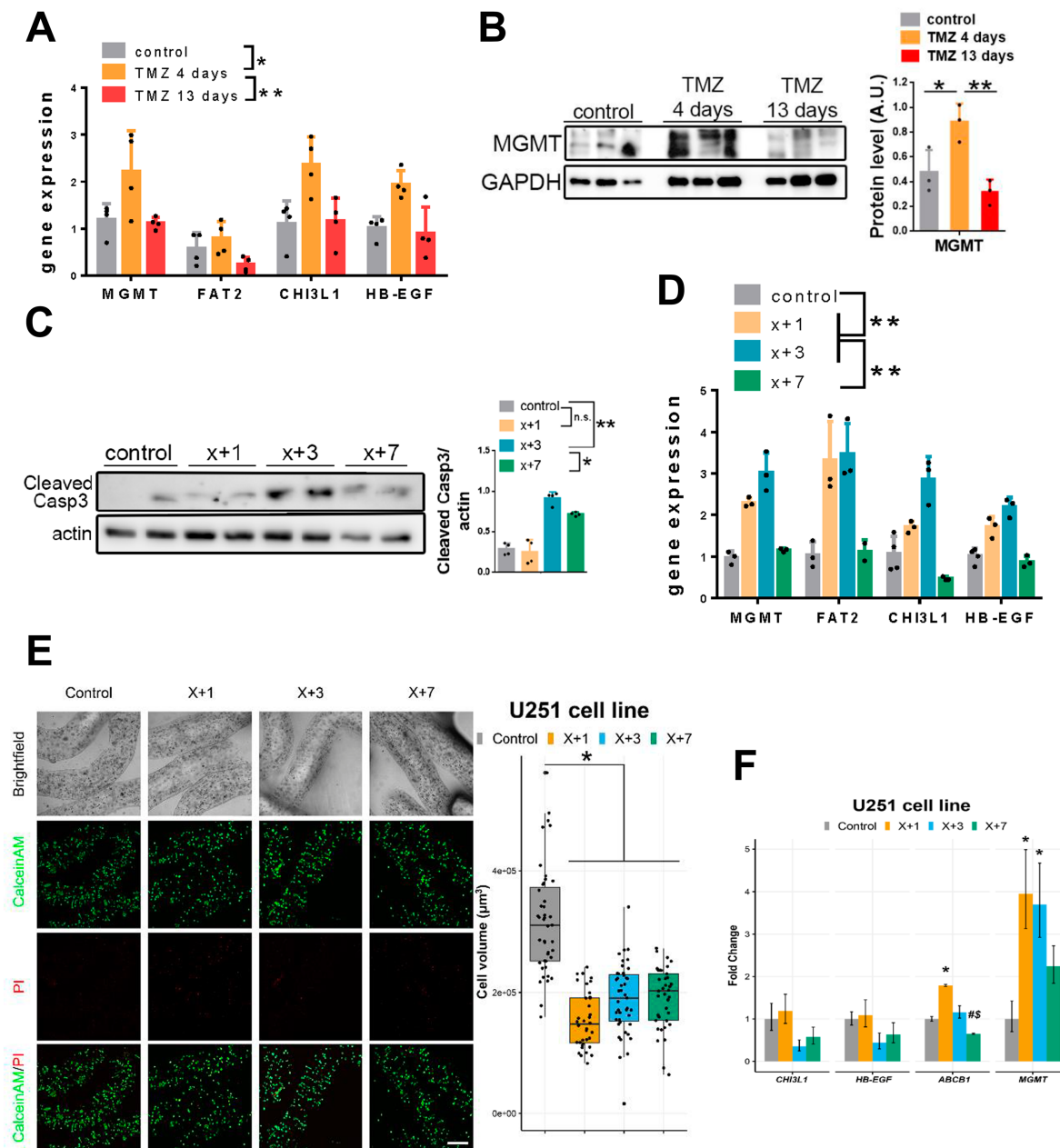
456

457 **Supplementary Fig. 1** Effect of increasing spacing between TMZ doses in SVZ
 458 EGFR wt/amp tumors. (A) Representative histological images stained with
 459 hematoxylin and eosin (H&E) from coronal brain sections of SVZ EGFR wt/amp
 460 tumors from (Fig. 2) of control, TMZ 4 days and TMZ 13 days treatment condition.
 461 Quantification of tumor area percentage is shown on the right (n=3). (B)
 462 Representative pictures of immunohistochemical (IHC) staining of ki67 in the
 463 same tumors from (A) and quantification of the percentage of proliferative cells
 464 on the right (n=3). (C) Representative pictures of IHC staining of Active Caspase3
 465 in SVZ EGFR wt/amp in the same tumors from (A). Quantification is shown on
 466 the right (n=3). * $P \leq 0.05$, n.s.=non significant.

467

468 As previously mentioned, persister cells represent an intermediate phenotype
469 arising before the development of TMZ resistance in gliomas²⁹. To test whether
470 different modalities of protracted TMZ treatment could affect the appearance of
471 these population of cells, the expression of persister genes was analyzed.
472 Notably, the expression of these markers was induced in tumors that had been
473 treated with the X+4, but not with the X+13, regime, as compared to untreated
474 animals (Figure 3A). Interestingly, one of these genes is *MGMT*, whose
475 expression is strongly associated with TMZ resistance⁴². It was confirmed that
476 *MGMT* protein was indeed being accumulated in the X+4 but not in the X+13
477 scheme (Figure 3B). These results suggest that extending the rest periods
478 between TMZ treatments not only improved the anti-tumor effect of the drug, but
479 also reduced the appearance of a persister state in the glioma cells.

480 In order to test whether the effect of protracted TMZ was cell-autonomous, SVZ-
481 EGFRwt cells were treated with different schedules of TMZ *in vitro*. An increase
482 was observed in caspase 3 cleavage in the X+3 scheme (Figure 3C),
483 accompanied by the accumulation of the expression of persister genes, including
484 *MGMT* (Figure 3D). Notably, an extended period between TMZ doses (X+7
485 scheme) did not produce a significant increase in Caspase 3 (Figure 3C) or in the
486 expression of persister genes (Figure 3D). These results suggest that the anti-
487 tumor mechanism of protracted TMZ doses is the same *in vivo* and *in vitro* and
488 does not depend on the tumor microenvironment.



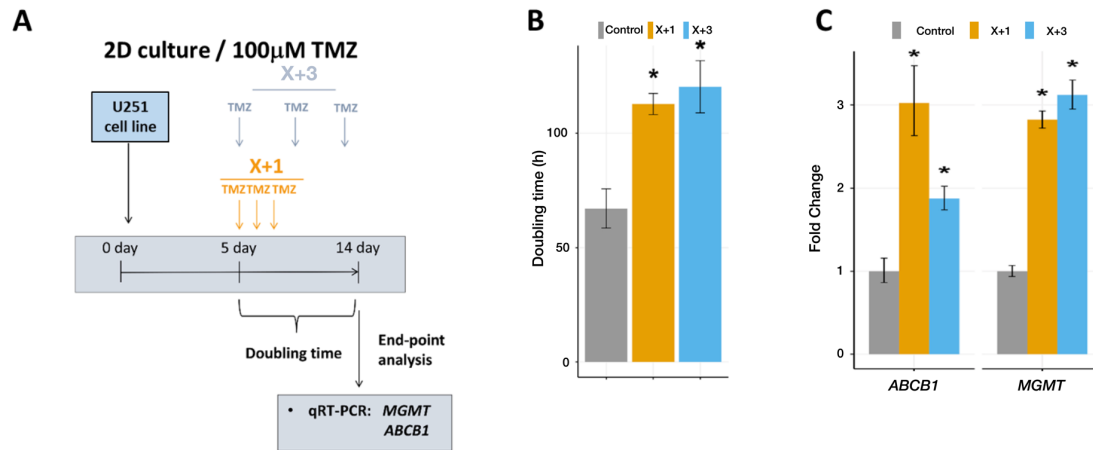
489

490 **Fig. 3** Increasing the spacing between TMZ treatments reduces the expression of
 491 persister-related genes *in vivo* and *in vitro*. (A) qRT-PCR analysis of persister-
 492 related genes in SVZ EGFR wt/amp tumors from (Fig. 2). *Actin* was used for
 493 normalization (n=4). (B) Western blot (WB) analysis and quantification of the
 494 expression of MGMT in SVZ tumors from (A). GAPDH was used for
 495 normalization. (C) WB analysis and quantification of Active Caspase3 in SVZ
 496 EGFR wt/amp cells treated *in vitro* with different TMZ schedules: control, TMZ 1
 497 day, TMZ 3 days and TMZ 7 days. *Actin* was used for normalization. (D) qRT-
 498 PCR analysis in SVZ EGFR wt/amp cells with the different treatment protocols.

499 *Actin* was used for normalization (n=4). (E) Representative confocal microscopy
500 images of U251 cells LIVE/DEAD labeled with CalceinAM/PI in alginate
501 microfibers after 28 days when the treatments and cultivation of cells were
502 completed. Quantification is shown as a box-plot on the right (n ≥ 3) Scale bar =
503 300 μm. (F) Relative gene expression levels of persister-related genes. *ACTB*
504 was used for normalization (n = 3). *P ≤ 0.05, **P ≤ 0.01, n.s.=non significant. #
505 indicates P<0.05 statistical difference compared to corresponding X+1
506 treatments. \$ indicates P<0.05 statistical difference compared to corresponding
507 X+3 treatments.

508

509 To confirm these results in human cells, U251 cells, grown in conventional 2D
510 conditions, were exposed to three doses of TMZ (100 μM), comparing daily (X+1)
511 and 3-day (X+3) schedules (Supplementary Figure 2A). Both regimes were able
512 to reduce the growth of the cells compared to the control, according to the change
513 in doubling time (Supplementary Figure 2B). However, in both cases an increase
514 in the expression of resistance-related markers in the TMZ treated cells was
515 detected (Supplementary Figure 2C). It was therefore determined to test the
516 effect of PTS in conditions where the cells would have slower growth. For this
517 purpose, U251 cells were immobilized in alginate microfibers and grown them in
518 3D conditions. The cells were then exposed to three doses of TMZ (100 μM),
519 comparing daily (X+1), 3-day (X+3) and 7-day (X+7) schedules (Figure 3E). In all
520 cases, TMZ reduced the growth of the cells (Figure 3F). Notably, X+7 decreased
521 the expression of persisters-related genes (*CHI3L1*, *HB-EGF*, and *ABCB1*) in
522 TMZ treated cells (Figure 3F). These results suggest that enlarging the intervals
523 between doses could reduce the appearance of a persister phenotype *in vitro* in
524 human glioma cells, at least in conditions of reduced proliferation.



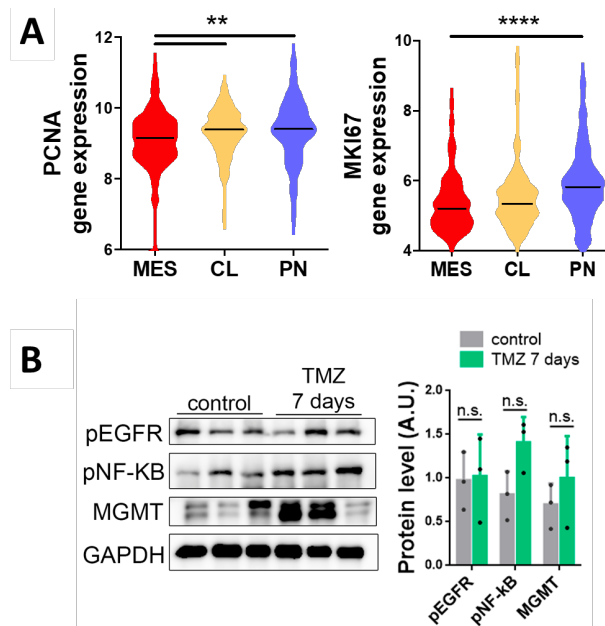
525

526 **Supplementary Fig. 2** Effect of TMZ *in vitro* in 2D cultures. (A) TMZ treatment
527 scheme in long-term 2D cell cultures of U251 glioblastoma cell line. (B) Doubling
528 Time (dt) calculated using RTCA 1.2.1 software illustrates the anti-proliferative
529 effects of TMZ treatments in U87 cell line. (C) Relative gene expression levels of
530 *ABCB1* and *MGMT* in U251 cells obtained by qRT-PCR. All values are expressed
531 as mean \pm SD and *ACTB* was used for normalization (n = 3).

532

533 **Protracted TMZ induced a change in the phenotype of slowly-proliferating**
534 **gliomas, mediated by epigenetic changes.**

535 Our previous characterization of the SVZ-EGFRwt model shows that these cells
536 express MES features^{30,31}. Interestingly, an *in-silico* analysis of two proliferation
537 markers, PCNA and MKI67, was performed, and it was found that MES gliomas
538 expressed the lowest levels of these genes (Supplementary Figure 3A),
539 suggesting that tumors with this phenotype are less proliferative than the other
540 two subtypes (classical (CL) and PN).



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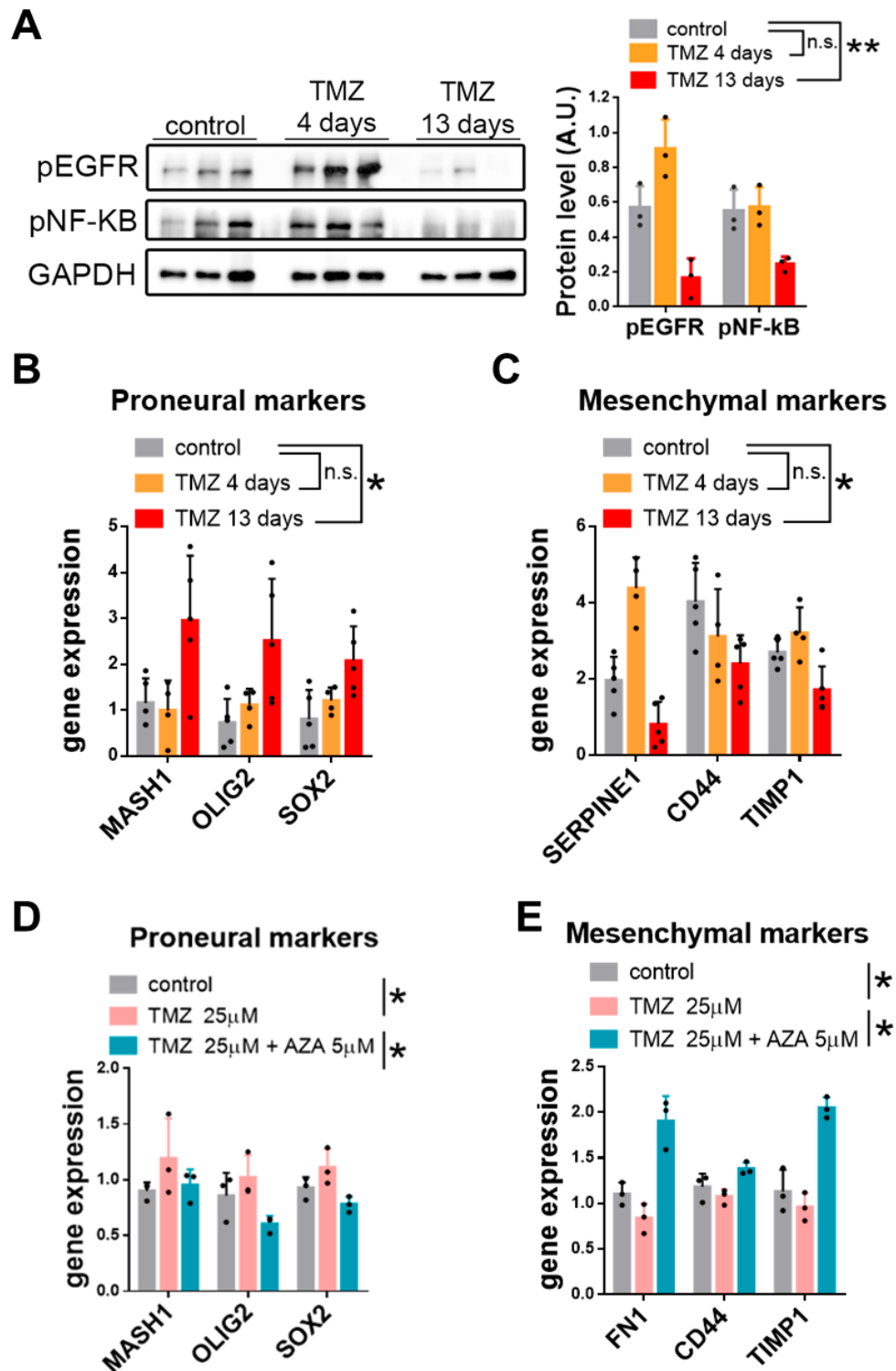
542 **Supplementary Fig. 3.** Expression of proliferation-related markers in different
543 subgroups of gliomas. (A) RNA-seq analysis of *PCNA* (left) and *MKI67* (right) in
544 a TCGA cohort stratified in to three groups: mesenchymal (MES), classical (CL)
545 and proneural (PN) tumors. (B) Western blot analysis of phosphorylated EGFR
546 (pEGFR) and NF-kB (p65) (pNF-kB) in SVZ-EGFRvIII tumors from Fig. 2.
547 GAPDH was used as a loading control. Quantification is shown on the right (n=
548 3).

549

550 It has previously been shown that the MES profile of SVZ-EGFRwt tumors
551 depends on the activation of the EGFR/NFkB signaling pathway. Notably, EGFR
552 has been associated with TMZ resistance in gliomas⁴³. The activation status of
553 this receptor in SVZ-EGFRwt tumors treated with TMZ was therefore explored,
554 both in responsive and non-responsive schedules. A clear downregulation of the
555 levels of phosphorylation of EGFR and NF-kB was observed in the X+13 tumors
556 (Figure 4A). In tumors from the X+4 regime there was an increase in phospho-
557 EGFR, although no significant changes were observed in the amount of phospho-
558 NFkB (Figure 4A), a MES driver in gliomas³⁹. The changes observed in the X+13
559 responsive tumors were paralleled by an increase in the expression of several

560 PN markers (Figure 4B) and the downregulation of the transcription of MES
561 genes (Figure 4C), compared to control tumors. These results suggest that less
562 intensive TMZ schedules might be reducing the aggressiveness of gliomas by
563 inducing a MES to PN phenotypic change. Notably, no changes in the amount of
564 phosphorylation of EGFR or NF- κ B were observed in TMZ-treated SVZ-EGFRvIII
565 tumors compared to controls (Supplementary Figure 3A), reinforcing the lack of
566 response of these gliomas to the drug.

567 To study the anti-tumor mechanisms of TMZ more deeply, SVZ-EGFRwt cells
568 were incubated *in vitro* in the presence of TMZ (25 μ M) for 8 days. As noticed in
569 tumors treated with TMZ, an increase in the expression of PN markers was
570 observed (Figure 4D). Notably, this effect was reverted in the presence of the
571 DNA-methyltransferase inhibitor 5-aza-2'deoxyctidine (AZA) (Figure 4D),
572 suggesting that TMZ might be inducing the expression of these genes by a shift
573 in the DNA methylation pattern, previously proposed as a mechanism of action
574 of this drug in glioma cells⁴⁴. A decrease in the expression of MES genes was
575 also noticed, which did not occur in the presence of AZA (Figure 4E). Therefore,
576 it was hypothesized that the anti-tumor effect of protracted schemes of TMZ might
577 be mediated, at least in part, by a MES-to-PN transition of the tumor cells induced
578 by epigenetic changes, the opposite of what would normally happen during tumor
579 progression⁴⁵. This would increase the effect of giving more time to persister cells
580 to revert their phenotypes to the PN phenotype.



581
582

583 **Fig. 4.** Mechanistic studies. Change of glioma phenotype for different TMZ
584 treatment schedules. (A) Western blot analysis and quantification of
585 phosphorylated EGFR (pEGFR) and NF-kB (p65) (pNF-kB) in SVZ EGFR wt/amp
586 tumors from (Fig. 2). GAPDH was used as a loading control (n= 3). (B-C) qRT-
587 PCR analysis of Proneural (B) and Mesenchymal subtype (C) related genes in

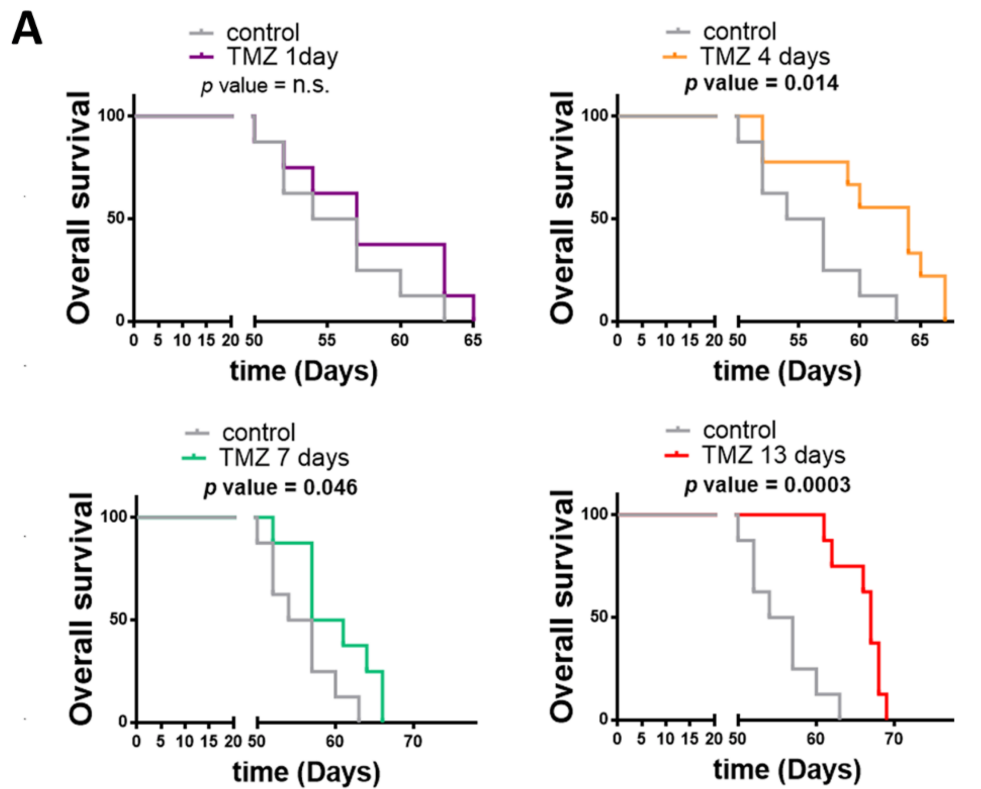
588 SVZ EGFR wt/amp tumors from (A). *Actin* was used for normalization (n=4). (D-
589 E) Analysis of the expression of Proneural (D) and Mesenchymal markers (E)
590 transcription by qRT-PCR in SVZ EGFR wt/amp cells cultured in the presence of
591 25 μ M TMZ, with or without azacytidine (AZA) (5 μ M). *Actin* was used for
592 normalization (n= 3). *P \leq 0.05, **P \leq 0.01, n.s.=non significant.

593

594

595 **TMZ dose can be increased in a protracted scheme to enhance the anti-**
596 **tumor effect and reduce toxicity.**

597 One of the potential benefits of long spacing between TMZ cycles could be a
598 reduction in the toxicity of the drug, which may perhaps allow the CT dose to be
599 increased. To test this hypothesis, the amount of TMZ administered to the
600 animals after the intracranial injection of SVZ-EGFRwt cells was increased to
601 50mg/Kg/day. The same schemes for TMZ treatment were used as in Figure 2,
602 and confirmed that the longest period between cycles was the most effective in
603 reducing tumor growth (Figure 5A). The day after the last TMZ cycle, blood was
604 collected from the animals to perform a white-cell count. One of the most common
605 adverse effects of chemotherapy with TMZ is myelosuppression, including
606 thrombocytopenia and leukopenia⁴⁶. Indeed, a reduction was observed in all the
607 numbers in the X+1 regime that reached a statistically significant value for the
608 decrease in thrombocytes (Figure 5B). Notably, extending the period between
609 doses was able to revert to normal the leukocyte and thrombocyte counts (Figure
610 5B), indicating that the toxicity of TMZ was reduced, even though the anti-tumor
611 effect was increased in comparison to the lower dose regime.



B

	TMZ (50mg/kg)			
	Untreated	1 day	7 days	13 days
<i>Hematology</i>				
Leukocytes ($\times 10^9/L$)	5,12 \pm 2,50	4 \pm 3,68	5,1 \pm 3,29	5,4 \pm 1,59
Neutrophiles ($\times 10^9/L$)	3,45 \pm 1,63	2,65 \pm 2,26	3,85 \pm 2,89	3,57 \pm 1,21
Lymphocytes ($\times 10^9/L$)	1,03 \pm 0,6	0,78 \pm 0,83	0,78 \pm 0,38	1,18 \pm 0,64
Trombocytes ($\times 10^9/L$)	426,75 \pm 172,57	131,25 \pm 91,94 (L)	339,5 \pm 283,6	349,25 \pm 195,22

612

613 **Fig. 5** Reduced toxicity of protracted TMZ schemes. (A) Kaplan-Meier overall
 614 survival curves of mice that were orthotopically injected with SVZ EGFR wt/amp
 615 cells and subsequently treated with intraperitoneal injections of a higher dose of
 616 TMZ (50 mg/kg per day) following the different TMZ protracted schemes (n = 9).
 617 (B) Hematology results in mouse blood samples from (A).

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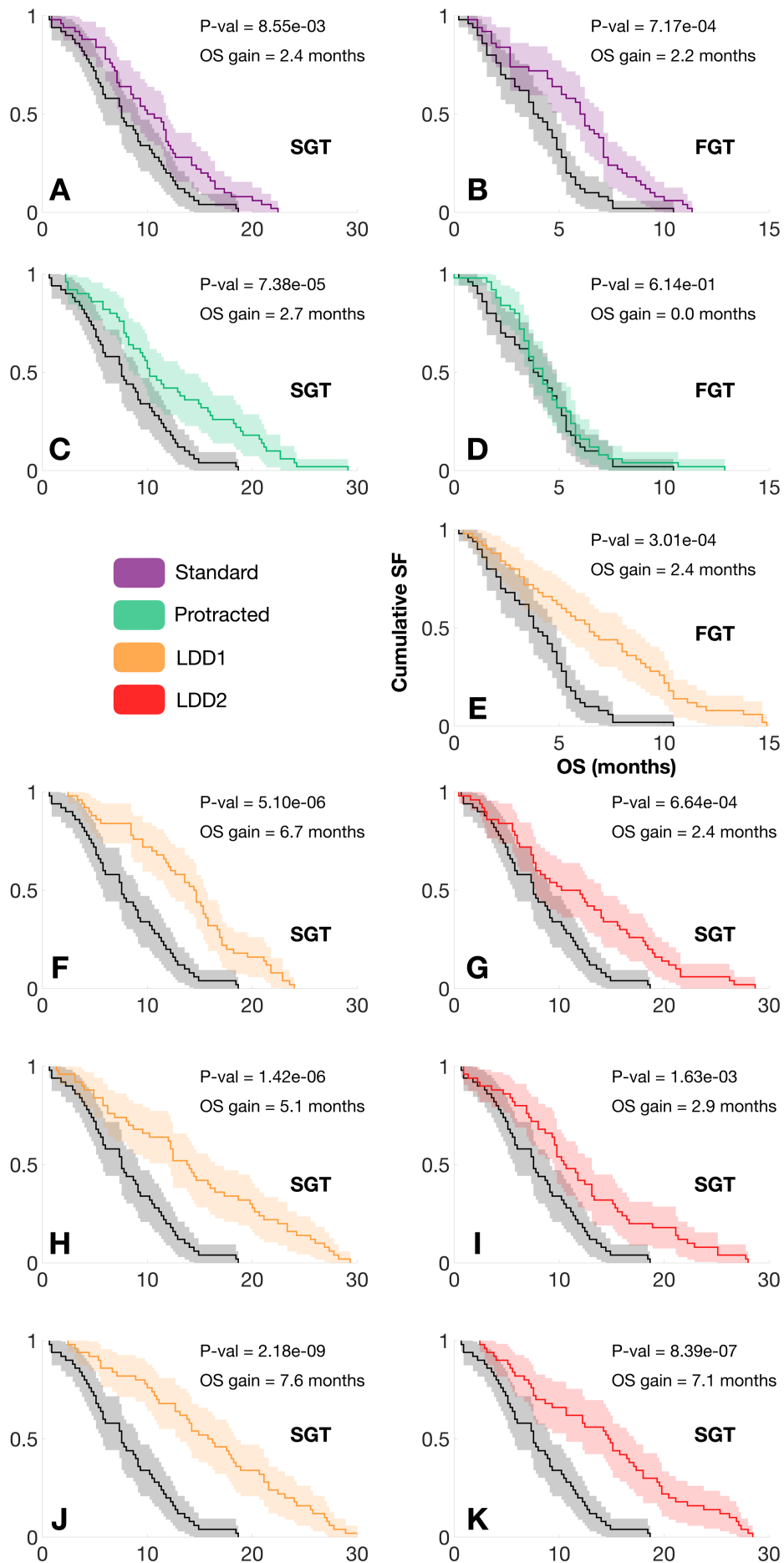
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624 **Virtual clinical trials suggest how to translate the results in experimental**
625 **models to human patients**

626 To extend previous *in vivo* findings, many sets of virtual clinical trials were
627 performed, based on the mathematical model with human parameters. The
628 potential effect of PTS on tumor growth dynamics was assessed by (i) enlarging
629 the rest periods between cycles, and (ii) testing long separation times between
630 individual doses without rest periods (Figure 6). Simulated tumors were
631 separated in slowly-growing and fast-growing GBMs. All *in-silico* patients were
632 given 30, 60 or 90 doses of TMZ 7 weeks after diagnosis (depending on therapy
633 regime).

634 Standard TMZ schemes (5 consecutive doses, rest period of 3 weeks) showed a
635 beneficial effect in terms of survival for both tumor types (Fig. 6A, 6B), with a
636 median survival difference of nearly 2 months, in line with clinical experience¹².
637 Increasing the rest period to 9 weeks improved survival for slow-growth tumors,
638 with an increased number of longer survivors (Fig. 6C). Fast-growing tumors did
639 not benefit from the increase in spacing between doses (Fig. 6D), in line with our
640 observations in animal models.

641 On the other hand, all trial branches benefited from increasing time intervals
642 between doses (without resting periods), both for slow and fast-growing tumors.
643 As the number of cycles given was increased, differences in median survival
644 increased remarkably.



646 **Fig. 6. Kaplan-Meier survival curves for PTS *in silico*.** Standard therapy (red)
647 consists of 6 cycles of TMZ, with 5 doses per cycle, and 23 days of resting period.
648 Protracted therapy (green) consisted of standard therapy, with a resting period of
649 9 weeks. Therapies with long spacing between doses 1 (LDD1, orange)
650 correspond to doses separated by 8 days without any rest periods. Therapies
651 with long spacing between doses 2 (LDD2, red) consisted of individual doses
652 spaced by 12 days without rest periods. **A)** Slow-growing simulated tumors with
653 6 cycles of standard therapy. **B)** Fast-growing simulated tumors with 6 cycles of
654 standard therapy. **C)** Slow-growing simulated tumors with 6 cycles of protracted
655 therapy. **D)** Fast-growing simulated tumors with 6 cycles of protracted therapy.
656 **E)** Fast-growing simulated tumors with 6 cycles of LDD1 therapy. **F)** Slow-
657 growing simulated tumors with 6 cycles of LDD1 therapy. **G)** Slow-growing
658 simulated tumors with 6 cycles of LDD2 therapy. **H)** Slow-growing simulated
659 tumors with 12 cycles of LDD1 therapy. **I)** Slow-growing simulated tumors with 18
660 cycles of LDD1 therapy. **J)** Slow-growing simulated tumors with 12 cycles of
661 LDD2 therapy. **K)** Slow-growing simulated tumors with 18 cycles of LDD2
662 therapy.

663

664 **DISCUSSION**

665 TMZ is the standard of care for newly diagnosed GBM, but the effect of this
666 alkylating agent is schedule-dependent⁴⁷. Genetic or acquired resistances to
667 TMZ can easily develop, and a strict regimen must be followed for a favorable
668 result to be obtained^{48,49}. The design of cytotoxic chemotherapy (CT) and
669 radiation therapy schedules is typically based on the basic principle of delivering
670 the maximum tolerated dose in the minimum time possible (MTDMT). The
671 rationale behind this is to avoid potential tumor repopulation during treatment, or
672 in the periods without treatment, and thus to achieve low tumor-cell numbers
673 compatible with patient cure. Although this is certainly the way to go when CT is
674 intended as a curative treatment, it is not obvious that it would be the best strategy
675 when it is known that treatment can only control disease for a limited time.

676

677 Intensification of TMZ delivery schemes has been studied for either newly
678 diagnosed³¹ or recurrent^{32,33} high-grade gliomas without positive results on OS
679 and with increased toxicity³⁸. However, no previous studies have considered
680 effective dose-reduction schemes with longer time spacing between treatments.
681 Different studies based on mathematical models have argued that cytotoxic
682 therapies with larger time intervals between doses could provide survival
683 benefits. However, these mathematical models are based on saturable growth
684 models, where tumors proliferate less on average as they grow larger, and the
685 phenomenon is lost when exponential growth models are considered²¹. The
686 situation would be worse in the context of super-exponential growth models²²
687 where the opposite result would be obtained, i.e. survival benefits from the
688 MTDMT approach.

689 This work studies, *in vitro*, *in vivo* and *in silico* using mathematical models, the
690 main factor limiting the effectiveness of treatment in gliomas: the development of
691 resistance. Resistance acquisition in the mathematical model was assumed to be
692 due to two biological facts: PN-MES transition, and persister induction by TMZ.
693 Accounting for resistances in our stochastic mesoscopic discrete simulation
694 model led to improved survival and reduced resistance when increasing the
695 interval between doses in virtual mice. The reduction of resistant cells stemmed
696 from the fact that longer spacing between doses allowed persister cells to revert
697 their phenotypes to PN. Clearly, when persistence time is longer than the spacing
698 between doses, persister cells receive additional TMZ doses and the emergence
699 of resistance is triggered. Very interestingly, for slowly-growing tumors, as TMZ
700 kills a fraction of both PN and MES cells, the resistance level at the end of the

701 simulation was observed to be smaller than its control counterpart, as the
702 spontaneous PN-MES transition is being reduced due to TMZ killing PN cells.
703 Thus, increasing the spacing between doses provides the same effect as a
704 reverse MES-PN transition. This setup was translated to the real world, both in *in*
705 *vitro* and *in vivo*. Results in mice were in accordance with model suggestions:
706 slow-growing tumors benefited most from PTS. In fact, fast-growing tumors did
707 not show an improvement in OS, as expected.

708 A phenotypic change in slowly-growing tumors subject to increased spacing
709 between doses was observed *in vivo*, with a reduction in the levels of
710 phosphorylated EGFR and NF- κ B, s associated with a MES-to-PN switch, as
711 recently shown³⁰. This transition could explain the reduced aggressiveness of the
712 tumors after long-cycle TMZ treatment, which does not seem to depend on
713 changes in proliferation and/or survival of tumor cells. Notably, our data suggest
714 that this MES to PN switch does not depend on the tumor microenvironment and
715 can be reverted in the presence of AZA, the known epigenetic regulator. Changes
716 in DNA methylation has already been associated with the response to TMZ in a
717 time and dose-dependent manner⁴⁴. Moreover, it has been previously shown that
718 the persister state is also linked with alterations in the levels of histone acetylation
719 and with chromatin remodeling processes²⁹. Therefore, it could be hypothesized
720 that epigenetic changes might be responsible for the appearance of resistances,
721 but also for some of the anti-tumor effects of TMZ, all linked to alterations in the
722 transcriptomic profiles of GBMs. Furthermore, our results might explain why
723 extensive TMZ treatment did not alter the survival of PN gliomas, but was
724 beneficial for the more aggressive MES subtype⁵⁰. Anyhow, these results not only
725 emphasize the potential clinical relevance of PTS for slowly-growing GBMs, but

726 also indicate that the biological assumptions taken in the model of action of this
727 compound are not far-fetched, and should be explored in deeper detail to keep
728 improving our knowledge about GBMs and the best way to treat them.

729 The experimental evidence shown here supports the fact that PTS could be
730 beneficial for GBM patients in terms of survival, resistance and toxicity so far. To
731 gain some insight into the improvements that could be expected in human
732 patients, many virtual trials were performed based on the mathematical model, to
733 explore the consequences PTS in clinical scenarios. The aim was to provide a
734 broad exploration of the outcomes of protracted regimes, and a proof of concept
735 that, if taken with caution, may be helpful in guiding the design of future clinical
736 trials.

737 The output yield by the virtual clinical trial was in agreement with the experimental
738 results obtained in this work. Standard TMZ therapy showed a moderate
739 improvement in terms of survival, in line with clinical experience. Fast-growing
740 tumors did not benefit from increasing the rest periods between cycles, but they
741 did benefit from enlarging the spacing between doses. Slow-growing tumors
742 benefitted not only from every alternative therapy scheme, but also from an
743 increase in the number of cycles given due to the reduction in the MES
744 component, and thus in tumor resistance. This points to alternative schemes that
745 would allow for more TMZ doses to be given, while keeping resistances stable
746 and with lower toxicity. There were several interesting implications from these
747 virtual trial studies. The first was that extending the rest period between 5-dose
748 cycles from 3 to 9 weeks showed a significant improvement in survival for patients
749 with slow-growing GBMs, suggesting an easy-to-apply upgrade in the standard
750 of care. The second was that there may be room for optimizing TMZ schedules

751 in GBMs, as the best improvements in survival came from schemes without rest
752 periods, and 8-12 days between doses. Due to their lower dose density, these
753 schemes could be less harmful in terms of toxicity and, following both virtual and
754 experimental results, a reduced resistance should also be expected, making
755 them an alternative option for clinical implementation.

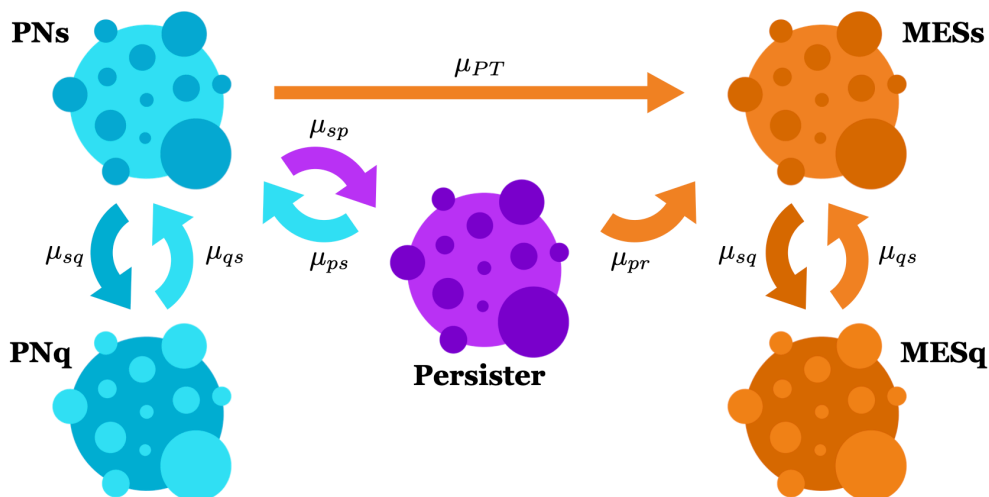
756 In conclusion, our combination of *in-silico* simulations, *in-vitro* and *in-vivo* studies
757 showed that TMZ administration schedules with increased time spacing between
758 doses may reduce toxicity, delay the appearance of resistances and lead to
759 survival benefits mediated by changes in the tumor phenotype, which was
760 especially important for slowly growing gliomas. The experimental results were
761 extended to human patients showing different ways to improve survival based on
762 the same concept.

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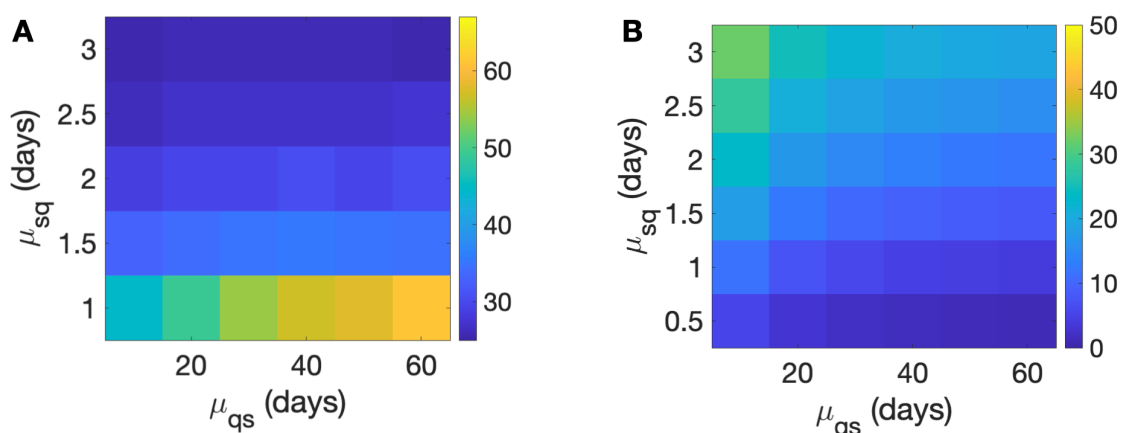
764 **ACKNOWLEDGMENTS**

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 777 analyzing results in human glioblastoma 2D and 3D cell culture.
 778



779
 780 **Supplementary Fig. S4.** Allowed cell transitions in the model. Both PN and MES
 781 cells share the same proliferative-quiescent dynamics. PNs can become MES
 782 either directly or due to TMZ exposure. In the latter case, the transition occurs
 783 through a transient reversible persister state.
 784



785

786 **Supplementary Fig. S5.** Overall survival in days **(A)** and Ki67 % **(B)** obtained

787 from different combinations of initial ranges of parameters μ_{sq} and μ_{qs} .

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