

Characterization of Temozolomide Resistance Using a Novel Acquired Resistance Model in Glioblastoma Cell Lines

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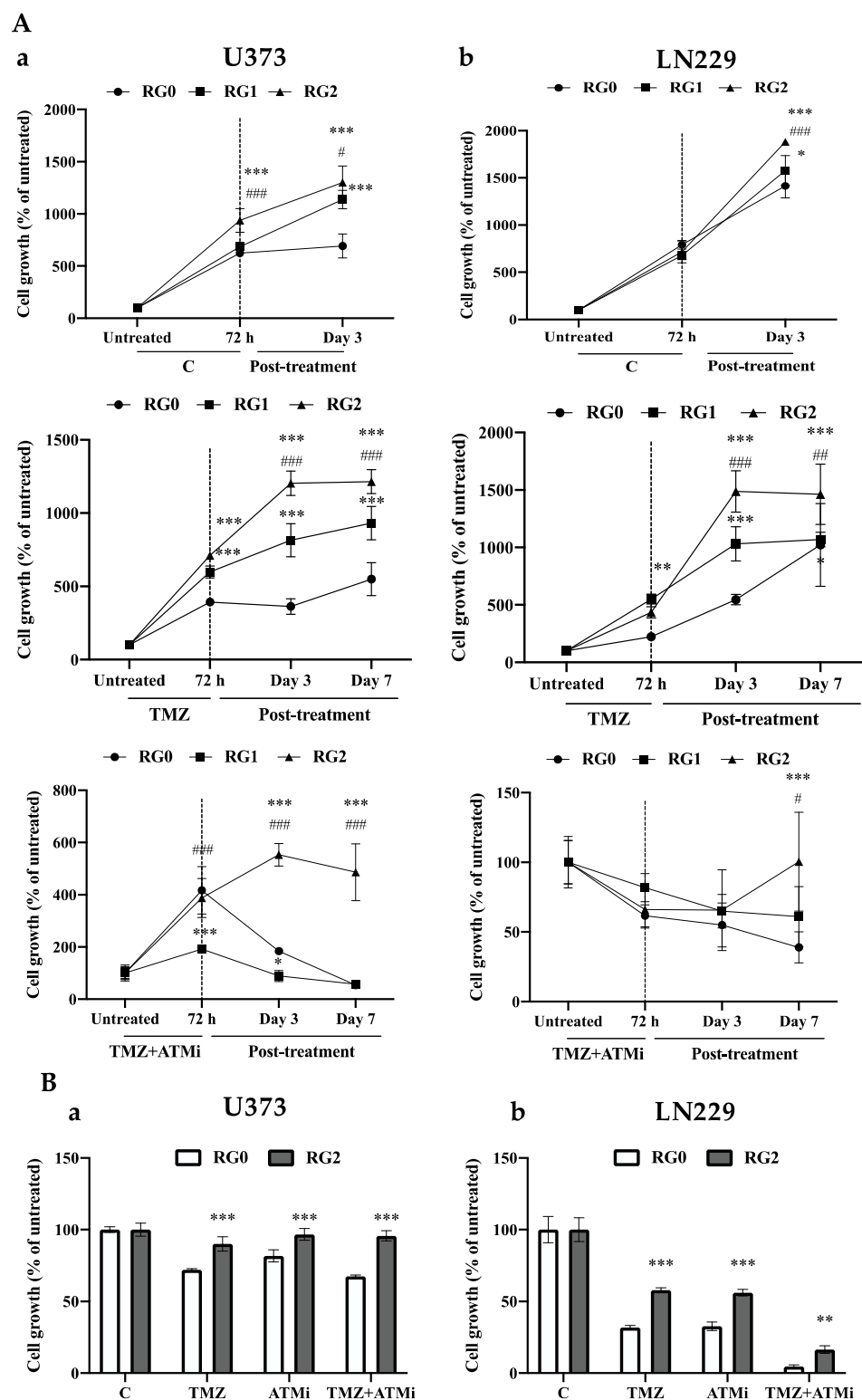


Figure S1. MTT assay in different passaged RG cells. (A) U373 RG cells (a) and LN229 RG cells (b) at passage 3 received the treatment with 0.2% DMSO as vehicle control (C), or TMZ (150 μ M), or TMZ+ATMi for 72 h. Thereafter, drugs and dead cells were removed by washing out with fresh medium. The viable cells were cultured for different periods (post-treatment) as indicated followed by MTT assay. (B) U373 RG cells (a) and LN229 RG cells (b) at passage 6 received the treatment with 0.2% DMSO as vehicle control (C) or TMZ (150 μ M) or ATMi (10 μ M) or TMZ+ATMi for 72 h followed by MTT assay. *, $p < 0.05$; **, $p < 0.01$; ***, $p < 0.001$, compared with RG0; #, $p < 0.05$; ##, $p < 0.01$; ###, $p < 0.001$, compared with RG1.

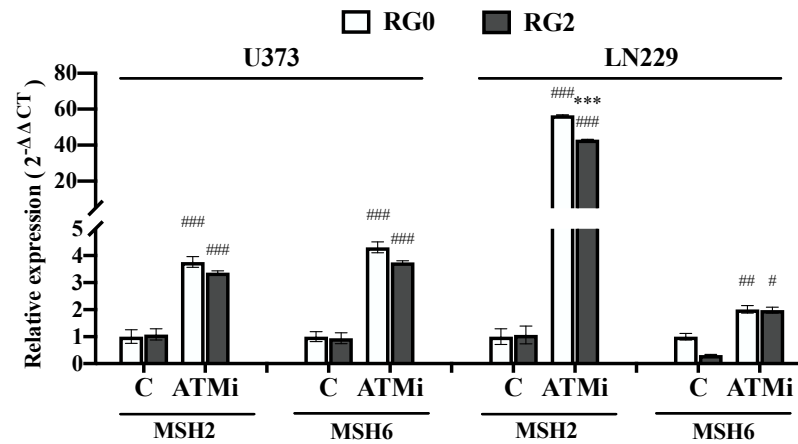


Figure S2. Expression of MMR genes in RG cells. Cells were treated with vehicle (0.2 % DMSO) or with ATMi (10 μ M) for 72 h. ***, $p < 0.001$, compared with RG0. #, $p < 0.05$, ##, $p < 0.01$; ###, $p < 0.001$, compared with a corresponding vehicle control (C).

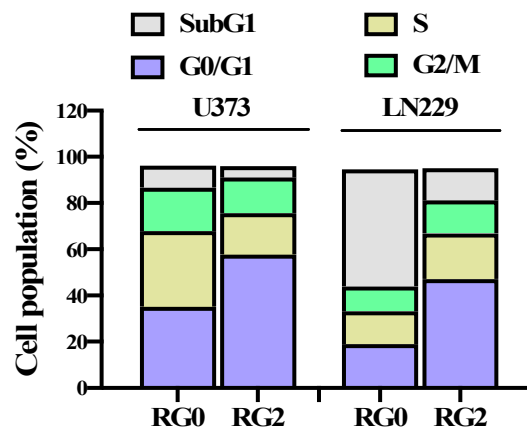
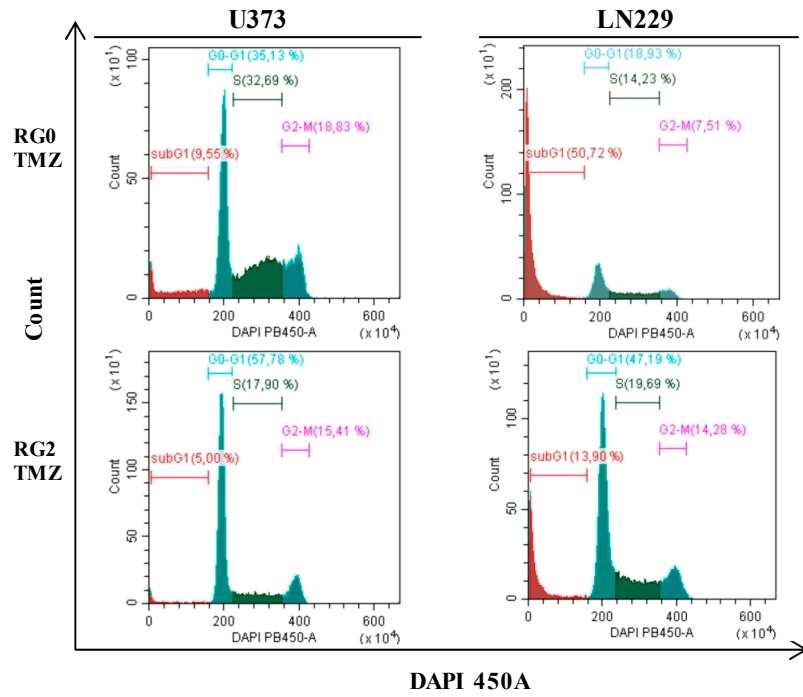


Figure S3. Flow cytometry-based cell cycle assay of RG cells including subG1 phase after 72 h TMZ treatment. In the subG1 phase, RG0 cells exhibited pronounced cell death compared to RG2 cells. Stacked bar graph summarizes the percentage of cells in the cell cycle phases including the subG1 in the experimental groups in the upper panel.

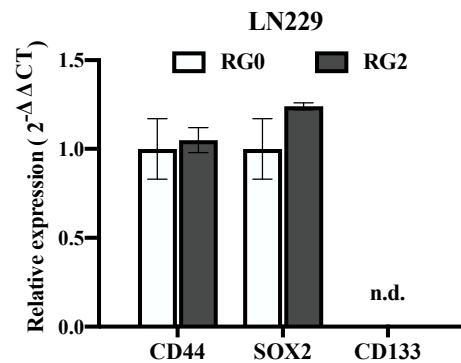
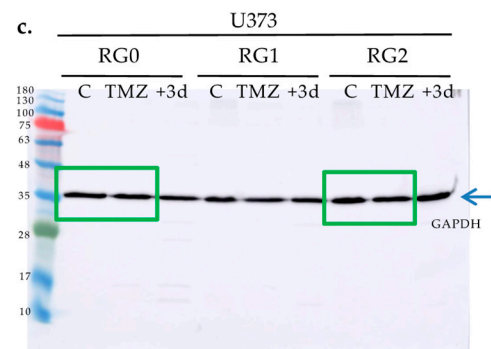
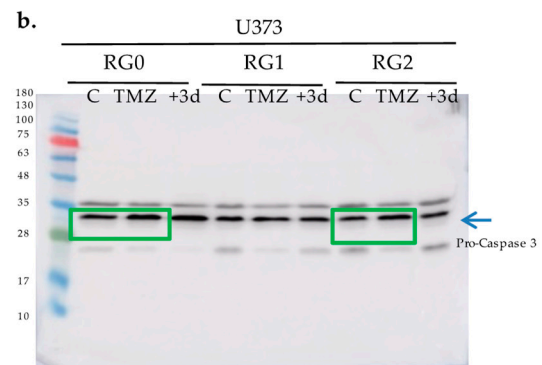
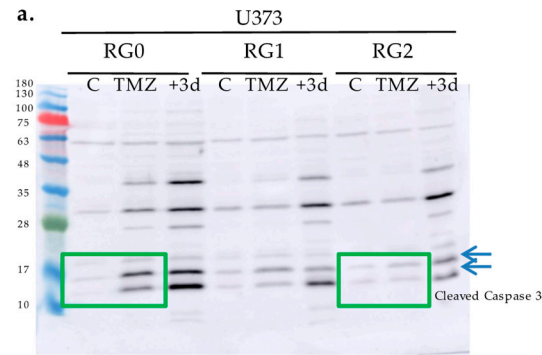


Figure S4. The expression of stem cell markers in LN229-RG cells detected by RT²-PCR. There was no significant difference in CD44 and SOX2 expression between RG0 and RG2. CD133 expression was not detectable (n.d.) in both RG0 and RG2.

A. U373



B. LN229

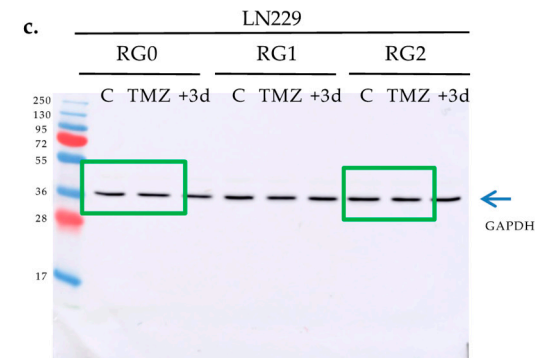
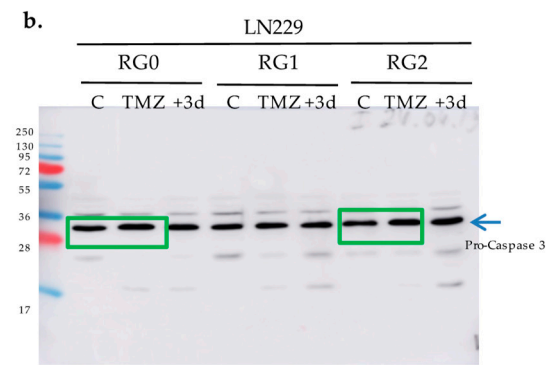
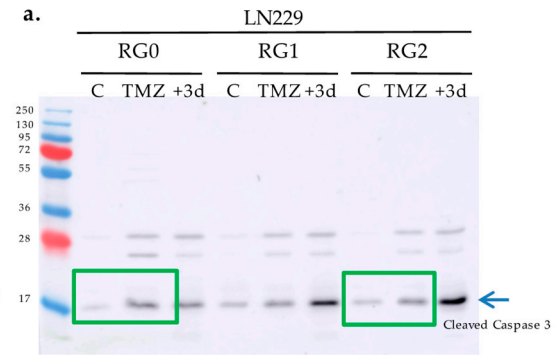


Figure S5. Original immunoblots for Fig. 6C. U373-RGs (A) and LN229-RGs (B) cells were collected 72 h after treatment with 150 μ M of TMZ or vehicle (C) for western blot. Another subset of regrown cells was harvested on day 3 of post treatment, whose data were not included in Fig. 6C. The blots shown in Fig. 6C for cleaved caspase 3 (a), pro-caspase 3 (b) and GAPDH (c) were marked in boxes.