Proteasomal Inhibition Selectively Kills Stem Cells in Glioma-Derived Cancer Cells Through Endoplasmic Reticulum-Associated Apoptosis

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b. Data
The data shown in figure should satisfy the following conditions:
- The data are accurate and consistent with the data that were actually obtained and submitted for publication.
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Each figure caption should contain the following information, for each panel where they are relevant:
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Statistics and general methods

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Identify the methodology applied for each protocol.

For a given study, answering the following questions will facilitate the identification:

- Was a simulation performed that simulated the data that resulted from all aspects of the AI system?
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G - Data Accessibility

Data deposition is mandatory for:

- Proteins, DNA and RNA sequences
- Macromolecular structures
- Crystallographic data for small molecules
- Functional genomics data
- Proteomics and molecular interactions

Deposition is strongly recommended for any datasets that are central and integral to the study; please consider the journal's data policy.

For datasets that are not deposited in a public repository, you should provide the accession codes for deposited data.

The data in GEOArchive files were deposited in the Gene Expression Omnibus (GEO) of NCBI (http://www.ncbi.nlm.nih.gov/geo/) under the accession number of GSE62356.

G - Dual use research

Could your study fall under dual use research restrictions? Please check biosecurity documents (see link list at top right) and list of select agents and toxins (APHIS/CDC) (see link list at top right). According to our biosecurity guidelines, provide a statement only if applicable.