Validated SF539 Xenograft Model: Subcutaneous Xenograft Tumor Model

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The Role of Xenografts in Studying Glioma Progression

Brain cancer, particularly malignant gliomas, represents one of the most aggressive and treatment-resistant forms of cancer, characterized by rapid progression, diffuse infiltration, and limited response to current therapies. Glioblastoma multiforme (GBM), the most prevalent and deadly form of brain cancer, continues to carry a poor prognosis, despite surgical resection, radiation, and chemotherapy. The biological complexity of brain tumors, including their pronounced heterogeneity, invasive growth patterns, and the restrictive nature of the blood-brain barrier, presents major obstacles to therapeutic development. *In vitro* models, while useful for mechanistic studies, fail to replicate the three-dimensional structure, cellular diversity, and microenvironment of brain tumors, reducing their predictive value in drug development. To address these limitations, xenograft models have become an essential platform in brain cancer research, enabling the study of tumor growth, invasion, and treatment response within a living organism. Cell line-derived xenografts (CDX) offer consistency and ease of use, while patient-derived xenografts (PDX) better preserve the molecular and histological characteristics of human tumors.

SF539 Cell Line

The SF539 cell line, originally derived from a recurrent glioblastoma multiforme (GBM) in a 34-year-old female, has been widely used in brain tumor research due to its distinct histological and molecular characteristics. Initially classified as a gliosarcoma, SF539 exhibits both gliomatous and sarcomatous features, with long-term cultures displaying elevated expression of extracellular matrix proteins such as collagen IV, fibronectin, and laminin. Notably, SF539 cells demonstrate high levels of low-density lipoprotein receptor-related protein (LRP), indicating enhanced lipid metabolism commonly associated with aggressive tumor behavior. Therapeutically, SF539 has shown significant sensitivity to temozolomide (TMZ), a frontline chemotherapeutic agent for GBM, which correlates with a highly methylated MGMT promoter and low MGMT protein expression, both of which are strongly associated with TMZ responsiveness. However, despite its frequent use, critical gaps remain, particularly in the comprehensive genomic characterization of SF539. Unlike other glioma models that have been thoroughly profiled for mutations in TP53, PTEN, and EGFR, SF539 lacks detailed mutational and transcriptomic data. In addition, its responses to novel or combinatorial therapeutic strategies have not been systematically studied, limiting its broader applicability in predictive modeling and translational oncology.

Altogen Labs Validated SF539 Xenograft Model

The SF539 glioma xenograft model, as established by Altogen Labs, involves the subcutaneous implantation of SF539 cells into immunocompromised mice to evaluate tumor growth and therapeutic response. SF539 cells are maintained under exponential growth conditions and prepared for inoculation at a density of one million cells suspended in 100 to 150 microliters of Matrigel. This suspension is injected into the flank of a hind leg, and tumor formation is monitored by palpation three times weekly. Once tumors reach an average volume of approximately 100 mm³, animals are randomized into treatment cohorts. Tumor volumes are measured daily using digital calipers, and mouse body weights are recorded twice weekly to monitor systemic toxicity. At study termination, animals are euthanized humanely, and tumors are excised, weighed, and photographed. Additional tissue samples are collected for downstream analyses, including preservation in RNAlater, snap freezing for molecular assays, or fixation for histopathological examination.





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Altogen Labs specializes in the development and implementation of xenograft-based in vivo models to assess the efficacy of investigational cancer therapies. These models, including the SF539 glioma model, provide a physiologically relevant system in which tumor cells are engrafted into immunodeficient mice using either subcutaneous or orthotopic methods. Xenograft studies at Altogen Labs are designed to replicate clinical tumor progression and therapeutic response, forming a critical component of preclinical drug evaluation. All FDAapproved anticancer agents have undergone assessment in similar in vivo systems. The comprehensive workflow includes selection of appropriate animal models, tumor cell line validation, compound dosing regimens, and rigorous data collection focused on tumor growth kinetics, histological features, and expression analyses of key molecular targets at both the mRNA and protein levels.



Figure 2. Tumor weight of SF539 cells in control, non-treatment mice and verotoxin treated mice at end of the study (Altogen Labs).

Subcutaneous SF539 Xenografts in Preclinical Glioma Research

Subcutaneous xenograft transplantation is a widely utilized method in preclinical cancer research, valued for its technical simplicity, reproducibility, and capacity for real-time monitoring of tumor growth. In the case of the SF539 gliosarcoma cell line, subcutaneous implantation into immunocompromised mice has proven effective in establishing consistent and measurable tumor growth, making it a valuable model for evaluating therapeutic efficacy *in vivo*. Originating from a recurrent glioblastoma, SF539 cells exhibit distinct biological characteristics, including a mesenchymal phenotype and sensitivity to DNA-damaging agents such as temozolomide. Subcutaneous xenografts derived from SF539 cells offer a reliable system for studying drug response and tumor progression, particularly in early-stage screening of novel chemotherapeutics. Although this model lacks the native brain microenvironment and associated tumor–stroma interactions, it remains highly informative for assessing tumor kinetics and pharmacological response under controlled conditions. Recent studies have employed SF539 xenografts to explore the effects of targeted therapies, DNA repair inhibitors, and combination regimens, contributing to a growing body of literature that supports the model's utility in translational neuro-oncology. Ongoing advancements in imaging technologies, molecular profiling, and co-engraftment strategies continue to improve the physiological relevance of subcutaneous xenografts, further reinforcing their role as a cornerstone of glioma drug development.

Case Study: FKBP9 Drives Glioblastoma Progression and Stress Resistance

In a study published by the *Journal of Experimental & Clinical Cancer Research*, Xu H., *et al.* investigate the role of FKBP9 in glioblastoma (GBM) progression and resistance to endoplasmic reticulum (ER) stress. The authors demonstrate that FKBP9 is highly expressed in high-grade gliomas and correlates with poor patient prognosis across multiple datasets. Functional studies using GBM cell lines such as SF539, LN-229, and T98G revealed that FKBP9 knockdown significantly suppresses proliferation, invasion, and stemness, while also downregulating key survival and stem cell markers including Bcl-2, XIAP, Oct4, and Sox2. *In vivo*, FKBP9 depletion reduced tumor burden in both chick chorioallantoic membrane and mouse xenograft models. Mechanistically, the study identifies activation of the ASK1–p38MAPK pathway as a driver of FKBP9-mediated oncogenic activity. Additionally, FKBP9 suppresses the IRE1α-XBP1 branch of the unfolded protein response, conferring resistance to ER stress inducers such as tunicamycin and thapsigargin. FKBP9 itself is subject to proteasome-mediated degradation upon ER stress, with lysine 265 identified as a critical ubiquitination site.

The study's methodology is comprehensive, integrating molecular and cellular assays with transcriptomic analysis and *in vivo* tumor models. The use of the SF539 glioma cell line, which exhibits mesenchymal characteristics, enhances the translational relevance of the findings. While the results strongly support FKBP9's role as an oncogenic factor and regulator of ER stress tolerance, limitations include the use of immunodeficient animals, which precludes assessment of FKBP9's immunological functions, and the absence of patient-derived xenograft validation. The research nonetheless contributes significantly to the field by identifying FKBP9 as a potential biomarker and therapeutic target in GBM. The findings suggest that targeting FKBP9 or its downstream signaling via the p38MAPK pathway, in combination with ER stress inducers, could improve treatment outcomes. Further studies are warranted to explore FKBP9's interactome, its role across GBM subtypes, and its potential as a predictor of therapeutic response in clinical settings.

Pt2ad Shows Superior Cytotoxicity Over Cisplatin in SF539

A newly synthesized dinuclear organoplatinum(IV) complex stabilized by adenine, referred to as Pt2ad, has demonstrated potent cytotoxic activity in a subset of human cancer cell lines, with particularly striking results observed in the SF539 glioblastoma cell line. In vitro assays conducted using the sulforhodamine B method revealed that Pt2ad achieved a 50 percent growth inhibition (GI50) in SF539 at a concentration of 1.47 micromolar, total growth inhibition (TGI) at 2.76 micromolar, and a lethal concentration 50 (LC50) at just 5.20 micromolar. Notably, these values indicate a greater overall cytostatic and cytotoxic effect than either cisplatin or temozolomide, which are current standards in glioblastoma treatment but were shown to be ineffective against SF539 in this study. SF539, derived from a glioblastoma multiforme tumor, is known for its resistance to conventional chemotherapeutics, making this response to Pt2ad particularly significant. The selectivity of Pt2ad's cytotoxicity, as demonstrated by its lack of effect on five other CNS cell lines, suggests a cell-specific mechanism that warrants further mechanistic investigation. These findings also reveal a disconnect between GI50 and LC50 values when comparing Pt2ad and cisplatin, highlighting a compound that is less immediately growth-inhibitory than cisplatin but more effective at halting cell division and inducing death over time. The experimental methodology is rigorous, leveraging the National Cancer Institute's 60-cell line panel, but further studies are necessary to evaluate in vivo efficacy, biodistribution, and blood-brain barrier permeability. Early computational data suggest Pt2ad may penetrate the bloodbrain barrier, though its high molecular weight and low water solubility may limit bioavailability. Further research should prioritize understanding the molecular underpinnings of SF539 sensitivity, potential platinum-DNA interactions, and structure-activity relationships, as well as assessing Pt2ad in orthotopic or patient-derived xenograft glioma models. These results position SF539 as a key model for evaluating the therapeutic potential of novel platinum-based compounds targeting glioblastoma.

Additional Case Study: Osimertinib Induces Paraptosis via ER Stress in Glioblastoma

The study by Hu *et al.*, published in *Cell Death Discovery* journal, investigates the mechanism by which Osimertinib, a third-generation EGFR tyrosine kinase inhibitor, induces cell death in glioblastoma (GBM), with a particular focus on non-apoptotic pathways. Among the GBM cell lines used, SF539 plays a prominent role in confirming the unique cytotoxic effects of Osimertinib. The authors demonstrate that Osimertinib induces extensive cytoplasmic vacuolation and cell death in SF539, along with LN-229, U87MG, and LN-18 cells, through a mechanism independent of caspase activation and autophagy. This mode of death, identified as paraptosis, is strongly associated with endoplasmic reticulum (ER) stress and activation of the unfolded protein response, specifically the PERK–eIF2 α –CHOP signaling pathway. In SF539 cells, Osimertinib treatment markedly increased the expression of CHOP and GRP78, as well as the accumulation of polyubiquitinated proteins, supporting the role of ER stress as a key mediator of cell death. RNA sequencing of Osimertinib-treated SF539 cells further confirmed the upregulation of ER stress-associated genes, reinforcing the centrality of the paraptotic response in this model.

The authors employed a comprehensive methodological approach, including immunoblotting, electron microscopy, viability assays, and *in vivo* xenograft studies, to characterize the effects of Osimertinib. In SF539, both the genetic and pharmacological modulation of TRIP13, an AAA+ ATPase, significantly influenced the cell's susceptibility to Osimertinib. TRIP13 overexpression in SF539 cells reduced ER stress marker expression and cytoplasmic vacuolization, thereby diminishing sensitivity to Osimertinib, while TRIP13 knockdown had the opposite effect. Importantly, treatment with the AKT inhibitor MK-2206 reduced TRIP13 levels and restored ER stress responses in SF539, enhancing the cytotoxicity of Osimertinib *in vitro*. These findings not only highlight SF539 as a representative and responsive glioma model but also underscore its utility in elucidating paraptotic mechanisms of cell death. While the study is limited by its use of immunocompromised animal models and requires clinical validation of TRIP13 as a biomarker, it introduces a promising strategy to overcome therapeutic resistance in GBM.

Structural Optimization and Selective Cytotoxicity in Glioma Cell Models

Hybrid compounds that combine Ciminalum and thiazolidinone scaffolds have shown strong anticancer activity, with notable effectiveness against central nervous system tumors. Among tested glioma models, the SF539 cell line demonstrated exceptional sensitivity, with significant growth inhibition and cytotoxicity observed at submicromolar concentrations. This high level of responsiveness suggests that SF539 is a particularly useful model for evaluating glioblastoma therapies involving small molecules that disrupt critical survival mechanisms in tumor cells. These compounds also showed selectivity, exhibiting minimal toxicity toward normal human lymphocytes, which indicates a potentially favorable therapeutic index. Structural variations on the thiazolidinone ring, especially at positions 3 and 5, had a substantial impact on biological activity. Substituents with different electronic properties altered the potency of the compounds, highlighting key structure activity relationships that can guide further optimization. While the data point to strong cytotoxic effects, the exact mechanisms by which these compounds act in SF539 cells remain undefined. Future

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investigations should determine whether the observed effects are driven by apoptosis, oxidative stress, or other forms of cell death. Moreover, linking the molecular profile of SF539, including DNA repair capacity, drug resistance markers, and metabolic traits, to its heightened drug sensitivity may reveal new therapeutic targets. These findings help expand our understanding of rational drug design for glioblastoma and support the development of targeted treatments for highly resistant brain tumors.

Oncogene Characteristics

Analysis of the SF539 glioblastoma cell line reveals that one of its principal oncogenic drivers is the MYC proto-oncogene, which is implicated in the regulation of mitochondrial DNA metabolism and transcription through its downstream effect on TOP1MT, a mitochondrial topoisomerase. According to Hu et al., SF539, along with other central nervous system (CNS)derived tumor cell lines in the NCI-60 panel, exhibits low expression of TOP1MT. This mitochondrial enzyme plays a critical role in mtDNA relaxation and structural maintenance of mitochondrial nucleoids, and its transcription is directly regulated by MYC via E-box binding motifs in the TOP1MT promoter and first intron. The functional consequence of MYCdriven TOP1MT expression is the coordination of nuclear and mitochondrial gene expression to support bioenergetic demands and cell proliferation. However, in SF539 cells, both MYC and TOP1MT are expressed at relatively low levels compared to other cancer types, which may suggest a divergent metabolic profile or altered mitochondrial regulation in glioblastoma cells relative to more MYC-driven malignancies such as leukemias. Despite this low basal expression, the regulatory axis involving MYC and TOP1MT remains biologically relevant. MYC amplification or overexpression has been shown to increase TOP1MT levels and promote mitochondrial biogenesis, thereby enhancing oxidative phosphorylation and supporting rapid tumor growth. The absence of such amplification in SF539 suggests a potential vulnerability that could be exploited therapeutically, particularly in strategies targeting mitochondrial function or energy metabolism. Moreover, given MYC's well-established role in driving cell cycle progression and evading apoptosis, its involvement in the modulation of mitochondrial gene networks positions it as a key integrator of metabolic and proliferative signals in glioblastoma. The unique oncogenic landscape of SF539, characterized by altered mitochondrial transcriptional regulation, supports the need for further research into how MYC and mitochondrial gene co-regulation influence glioma progression and therapeutic response. Future studies should investigate whether manipulation of the MYC-TOP1MT axis can sensitize SF539 cells to mitochondrial stress or targeted inhibitors, thereby opening new avenues for glioblastoma treatment.

Altogen Labs is a leading preclinical contract research organization (CRO) offering a comprehensive suite of biology services desianed to support oncology drua development and translational research. As a specialized provider of in vivo and in vitro pharmacology services, Altogen Labs conducts pharmacology and toxicology testing, IC50 determination across more than 100 human cancer cell lines, and anti-tumor efficacy studies using over 90 validated cell line-derived xenograft (CDX) models and over 30 patientderived xenograft (PDX) models. The laboratory also offers custom liposome encapsulation for siRNA, mRNA, and DNA, ELISA and cell-based assay development, and generation of stable cell lines for long-term gene expression or knockdown studies. Additional services include in vivo toxicology studies in mice and rats (including LD50 and teratoma formation assays), RNA interference (RNAi) technologies such as inducible knockdown models and in vivo siRNA delivery, as well as GLP-compliant cell banking and cryopreservation.



Figure 3. Available *in vivo* xenograft services at Altogen Labs for SF539.

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All research conducted at Altogen Labs adheres to IACUC and GLP regulations, ensuring compliance with preclinical safety and efficacy standards for regulatory submissions. Among Altogen Labs' in-house validated xenograft models is the SF539 glioblastoma xenograft, a robust and widely used model for studying gliosarcoma biology and therapeutic response. The SF539 cell line, derived from a recurrent glioblastoma multiforme tumor of a 34-year-old patient, is notable for its expression of extracellular matrix proteins such as collagen type IV, laminin, and fibronectin, and is part of the NCI-60 cancer cell panel. Altogen Labs utilizes the SF539 model to evaluate both monotherapies and combination treatments, including temozolomide and multi-kinase inhibitors like PHA-848125, which has demonstrated brain-penetrant activity. In this model, SF539 cells are subcutaneously implanted in immunocompromised mice, and tumor growth is monitored until reaching 100-150 mm³ before treatment begins. Altogen provides end-to-end studv Labs



Figure 4. Altogen Labs provides *in vivo* toxicology services to assess dose response, safety, and biological activity for preclinical drug development.

execution, including tumor measurement, body weight tracking, tissue collection, necropsy, histopathology, and downstream molecular analysis such as RNA and protein extraction, RT-PCR, and immunohistochemistry. Additional study customization options include variable dosing schedules, delivery routes (e.g., IV, IP, intratumoral, oral), imaging studies, and blood chemistry analysis. The SF539 xenograft model at Altogen Labs serves as a vital preclinical platform for evaluating the efficacy and safety of experimental brain cancer therapies, offering clients detailed reporting that includes raw data, statistical analysis, and interpretive discussion.

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