

Potent Targeting of the STAT3 Protein in Brain Cancer Stem Cells: A Promising Route for Treating Glioblastoma

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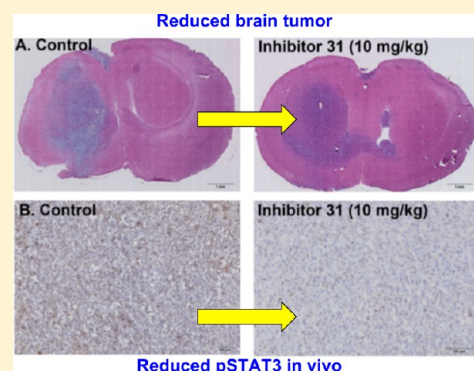
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Supporting Information

ABSTRACT: The *STAT3* gene is abnormally active in glioblastoma (GBM) and is a critically important mediator of tumor growth and therapeutic resistance in GBM. Thus, for poorly treated brain cancers such as gliomas, astrocytomas, and glioblastomas, which harbor constitutively activated *STAT3*, a *STAT3*-targeting therapeutic will be of significant importance. Herein, we report a most potent, small molecule, nonphosphorylated *STAT3* inhibitor, **31** (**SH-4-54**) that strongly binds to *STAT3* protein ($K_D = 300$ nM). Inhibitor **31** potently kills glioblastoma brain cancer stem cells (BTSCs) and effectively suppresses *STAT3* phosphorylation and its downstream transcriptional targets at low nM concentrations. Moreover, in vivo, **31** exhibited blood–brain barrier (BBB) permeability, potently suppressed glioma tumor growth, and inhibited p*STAT3* in vivo. This work, for the first time, demonstrates the power of *STAT3* inhibitors for the treatment of BTSCs and validates the therapeutic efficacy of a *STAT3* inhibitor for GBM clinical application.

KEYWORDS: *STAT3*, BTSCs, glioblastoma, small-molecule inhibitor, antitumor cell effects



Glioblastoma (GBM) is considered the most aggressive and lethal of brain cancers, with a median survival after treatment of approximately 15 months; shockingly, these modest results can only be achieved in the relatively young, otherwise healthy patients.¹ Moreover, GBM is neither preventable, nor detectable at a stage when early treatment might be more effective. Furthermore, despite intensive research, major improvements in overall survival have remained elusive.

Brain tumors have been demonstrated to contain rare subpopulations of brain tumor stem cells (BTSCs), which possess the cardinal stem cell properties of clonogenic self-renewal, multipotency, and tumorigenicity.² The extensive self-renewal and proliferative capacity of BTSCs coupled with their insensitivity to conventional radio- and chemotherapies^{3,4} suggest that they are integral to the growth and post-treatment recurrence of GBM. As such, BTSCs represent a reservoir of disease that require novel therapeutic approaches to eliminate in order to improve the outcome of GBM. Recently, Carro et al.⁵ demonstrated that the signal transducer and activator of transcription 3 (*STAT3*) gene is abnormally active in GBM,

and is a critically important mediator of tumor growth and therapeutic resistance in GBM. Poorly treated brain cancers such as gliomas, astrocytomas, and glioblastomas harbor constitutively activated *STAT3*. In addition, a growing body of recent evidence gathered using a variety of different small molecules that indirectly inhibit *STAT3* by targeting upstream molecules such as the JAK family members⁶ strongly suggest that *STAT3* signaling is crucial for the survival and proliferation of BTSCs and GBM both in vitro and in vivo.⁷ However, because of their broad targeting nature, these drugs have limited translational potential due to numerous side effects. Hence, drugs with the ability to more specifically block *STAT3* activity may provide effective treatment of GBM.

Briefly, *STAT3* is a member of the *STAT* family of transcription factor proteins. *STAT3* is activated through phosphorylation of tyrosine 705 (Y705) that initiates complexation of two phosphorylated *STAT3* monomers (p*STAT3*).

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66 pSTAT3 homodimers are mediated through reciprocal STAT3
 67 Src Homology 2 (SH2) domain-pY705 STAT3 interactions.
 68 pSTAT3:pSTAT3 homodimers translocate to the nucleus and
 69 bind DNA, promoting STAT3 target gene transcription.⁸
 70 Targeting STAT3 has been achieved with dominant negative
 71 constructs, oligonucleotides,⁹ or, most commonly, phospho-
 72 peptidic agents that mimic the native pY705 containing binding
 73 sequence^{10–13} or non-native STAT3-binding sequences.^{14,15}
 74 Unfortunately, these inhibitors, possessing pTyr-like groups, are
 75 poorly cell permeable and rapidly metabolized in vivo, which
 76 has limited their use in the clinic. To circumvent these
 77 problems, our group has developed novel small molecule
 78 STAT3 inhibitors not containing a phosphate ester.^{16,17} Using
 79 structure-based design, targeting the pTyr-SH2 domain
 80 interactions, salicylic acid-based inhibitors have been identified
 81 that block STAT3 dimerization and DNA-binding, namely, **BP-**
 82 **1-102** (1, Figure 1). **BP-1-102** potentially suppresses cell

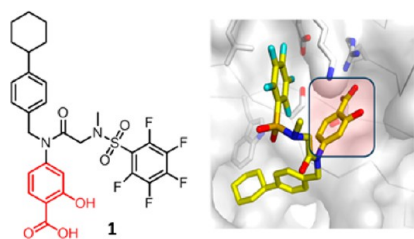


Figure 1. In silico docking images of inhibitor 1.

83 proliferation, migration, invasion, and motility. It showed little
 84 or no effect on phosphorylation of Jak-1/2, Shc, Src, or Akt.
 85 **BP-1-102** exhibited potent antitumor effects in in vivo
 86 xenograft models of lung and breast cancer. Western blots of
 87 tumors showed repression in pSTAT3 in a dose-dependent
 88 manner.^{18–20} We hypothesized that since the STAT3 pathway
 89 is a key oncogenic driver in brain tumor stem cells, direct
 90 inhibition of STAT3 would provide a targeted route for
 91 managing GBM. In this study, we have optimized 1 to be more
 92 potent, possess reduced pharmacokinetic liabilities and
 93 effectively penetrate the blood brain barrier (BBB).

94 A library of **BP-1-102** analogues possessing both prodrugs
 95 and potential bioisosteres, as well as salicylic acid mimics, were
 96 prepared (Table 1). For improving cell and BBB permeability,
 97 prodrugs were synthesized to conceal the carboxylate
 98 functionality. In this family, alkyl (21 and 22), acetoxymethyl
 99 (AOM) (23), pivaloyloxymethyl (POM) (24), and the
 100 acetylated prodrug (19) were prepared to enhance cell
 101 penetration. We prepared a nonhydrolyzable phosphonate,
 102 18, as well as its prodrug, 16. In addition, the relative ring
 103 positions of the hydroxy- and carboxylate groups of the salicylic
 104 acid were inverted (34), the hydroxyl substituent replaced with
 105 a fluoride (33) or removed entirely and replaced with a
 106 hydrogen atom (31). The fluorinated analogue, 33, was
 107 prepared to reduce the charge on the carboxylate and preclude
 108 phase II glucuronidation of the phenol. To investigate the
 109 salicylic acid's role in binding STAT3, an *N*-hydroxyl amino
 110 (32), sulfonamide (10–13), sulfonamine (14), tetrazole (29
 111 and 35) and *N*-hydroxy-oxamic acid (25) derivatives were
 112 prepared. As negative controls, we prepared 10, 11, and 30,
 113 which possessed hydrophobic naphthyl or benzene groups in
 114 place of the salicylic acid.

115 First, the library was screened for biological activity against
 116 GBM BTSC line 30M, derived from a GBM patient. Cell

Table 1. Focused Library of Inhibitors Consisting of Prodrugs and Bioisosteres of the Salicylic Acid Functionality

#	R ₁	#	R ₁	#	R ₁
8		18		28	
9		19		29	
10		21		30	
11		22		31	
12		23		32	
13		24		33	
14		25		34	
15		26		35	
16		27		36	

viability following drug treatment was assessed after 72 h using
 an alamarBlue assay.^{6,21} IC₅₀ values were derived and compared
 to **BP-1-102**. Consistent with previous findings, **BP-1-102**
 exhibited low micromolar activity (Figure 2A). More
 encouragingly, we identified a number of more potent
 inhibitors exhibiting low nanomolar IC₅₀ values. Lead
 compounds were then evaluated against a panel of BTSCs,
 25EF, 67EF, 73E, 84EF, and 127EF, which are molecularly
 heterogeneous human GBM BTSCs (Figure 2B). Compounds
 displayed IC₅₀ values ranging from 0.1 to 3.8 μM in comparison
 to **BP-1-102**, which displayed values above 2–5 μM (BTSC
 30M). Compounds SH-05-19 (19), SH-04-54 (31), SH-05-07
 (32), and SH-05-23 (33) showed potent activity, with IC₅₀s

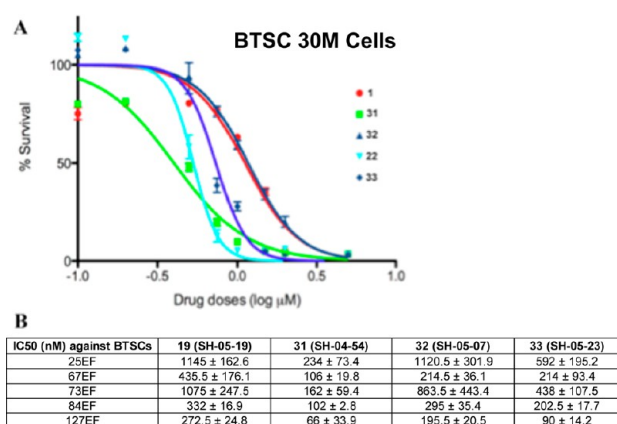


Figure 2. (A) Comparison of BP-1-102 (1) against the top four from the new library in BTSC30M; (B) IC₅₀ values of top four inhibitors against BTSCs 25EF, 67EF, 73E, 84EF, and 127EF.

STAT5 (Supporting Information), we cannot attribute the observed activity to anti-STAT5 activity.

Next, to examine phosphopeptide:STAT3 SH2 domain disruption, compounds were subjected to a fluorescence polarization (FP) assay.^{23,24} As expected, prodrugs 15–16 and 19–24 displayed no disruptive potency. Inhibitors 31, 32, and 33 showed good inhibitory potency, disrupting STAT3-phosphopeptide interactions with K_i s ranging from 10 to 30 μ M. Encouragingly, 33, 31, and 32 were selective for STAT3 over STAT1, a tumor suppressor protein (2-, 4-, and >10-fold selectivity for STAT3, respectively (Supporting Information). Concerned that binding affinity was due to hydrophobic aggregation, we tested 30, which lacks the salicylic acid, and found that it had no binding affinity for the STAT3 protein (Supporting Information), thus confirming the requirement for the carboxylate for 31's binding potency.

Next, 19-, 31-, and 32-mediated inhibition of cellular levels of activated pSTAT3 activity were determined using Western Blot analysis for phosphorylated STAT3 (pY705 and pS727) as well as for downstream target genes, Bcl-xL and Cyclin D1 (Figure 4A). Inhibitors exhibited concentration-dependent and potent

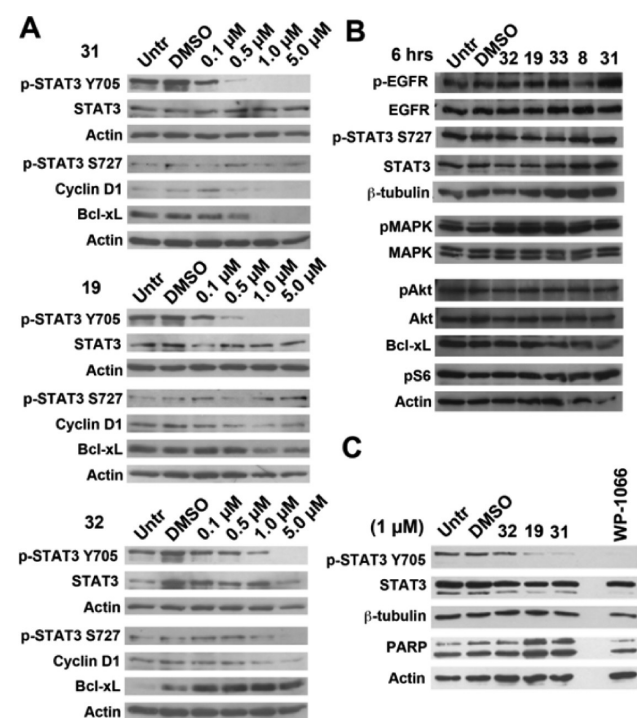


Figure 4. (A) Compounds effectively attenuate pSTAT3 Y705 with no effect on pSTAT3 Ser727; (B) exhibit no off-target effects in various signaling pathways; (C) Western blot analysis of top agents in comparison against Jak2 inhibitor, WP1066.

(nM) suppression of pSTAT3 (Y705) with no effect on the total STAT3 levels or on pSTAT3 (S727). Of note, blockade of the STAT3 SH2 domain should not inhibit S727 phosphorylation, suggesting that our molecules were binding the SH2 domain. Further Western analysis for inhibition of upstream kinase targets including, Jak2, Src, (Supporting Information), Akt, MAPK, and EGFR showed no off-target effects (Figure 4B). Most encouragingly, dose-dependent decreases in pSTAT3 levels were observed as well as potent inhibition of downstream targets involved in cell growth and survival (Cyclin D1 and Bcl-xL). Notably, 31 exhibited concentration-depend-

Figure 3. (A) Top three hits showed minimal cytotoxicity against normal human fetal astrocyte cells; (B) SPR curves displaying the binding affinity of the best hit, 31, to STAT 3/5 proteins. Studies were done at multiple concentrations (5.000, 1.667, 0.556, and 0.185 μ M).

Next, to determine kinetic association and dissociation rate constants (k_{on} and k_{off}), surface plasmon resonance (SPR) binding experiments were performed using a ProteOn XPR36 (Biorad) with full length His-tagged STAT3 protein immobilized on THE sensor chips. Promisingly, lead agents exhibited nanomolar binding potencies with 31, the most potent, exhibiting a K_D (k_{off}/k_{on}) of 300 ± 27 nM (Figure 3B). Thus, 31 represents a most potent, nonphosphorylated, small molecule STAT3 inhibitor. While phosphopeptides have been reported to bind to STAT3 with nanomolar potency, these agents have not shown in vivo efficacy.²² Comparative SPR analysis against STAT5, an analogous member of the STAT family proteins, showed modest selectivity ($K_D = 464$ nM). Since the BTSCs evaluated do not harbor hyperactivated

ent decreases in pSTAT3 levels that correlated well with observed cytotoxicity and downstream target suppression. Treatment with **31** (500 nM) silenced pSTAT3 signaling in 147EF cells. Inhibitors **19** and **32**, while less potent, exhibited nanomolar inhibition of pSTAT3.

To evaluate potential off-target effects, **19**, **31**, and **32** were screened in vitro for activity against cancer related kinases, c-Src, ERK1, Akt, Jak1, and Jak2 at 5 μ M (Supporting Information). Inhibitors exhibited moderate to negligible activity against the kinases tested. Compounds showed negligible inhibition of Akt1, Erk1, and Jak1, while only exhibiting modest inhibition of Jak2, c-Src, and Jak2 (~50% inhibition at 5 μ M). Since the concentration of inhibitor required to elicit effective kinase inhibition was 15-fold higher than the IC₅₀ values in BTSCs, we concluded that activity was not a result of Jak kinase inhibition.

To comprehensively investigate potential off-target effects, **31** was subjected to a kinome screen (101 diverse kinases, DiscoverX) as well as a protein and receptor screen (21 biologically important G protein-coupled receptors (GPCRs)). For kinome screening, ultrasensitive quantitative PCR (qPCR) was used to measure levels of immobilized kinases after treatment with **31**.¹² The GPCR screening employed the PathHunter β -arrestin assay platform (DiscoverX) to evaluate **31**. Encouragingly, **31** showed no off-target activity against any of the 21 GPCRs tested (500 nM). Moreover, in the kinome screen (500 nM), **31** showed negligible effects against SH2 (Jak1/2) and SH3 (Fes, Fer, and Fyn) containing kinases with the exception of 35% inhibition of PDGFRB, a kinase implicated in glioblastoma.²⁵ Concerned that **31** may act on PDGFRB to inhibit STAT3 activity, we conducted Western Blot analysis for activated PDGFRB in BTSCs evaluated as well as to determine the IC₅₀ value of **31** against PDGFRB in vitro. Encouragingly, we found that PDGFRB activity was not present in any of the BTSCs examined (Supporting Information), and furthermore, as assessed by an enzymatic assay, **31** did not inhibit PDGFRB at concentrations of 10 μ M (Supporting Information). Thus, to the best of our knowledge **31** directly inhibits STAT3.

Next, we compared **31** to a potent Jak2 inhibitor, WP1066, in 73 M BTSCs (Figure 4C).⁶ Western blot analyses showed equipotent suppression of pSTAT3 at 1 μ M. In a cytotoxicity assay, **31** was found to be more potent (30 M, **31** IC₅₀ = 0.43 μ M, cf. WP1066 IC₅₀ = 1.8 μ M, and in 73 M, **31** IC₅₀ = 1.03 μ M, cf. WP1066 IC₅₀ = 2.1 μ M (Supporting Information). Notably, **31** does not inhibit Jak2 at therapeutic doses and appears to function through direct STAT3 inhibition.^{26,27} In addition, Western blot analysis of **31** vs **1**, where comparative effects against pSTAT3 and pAkt were measured, as well as induced apoptosis as measured by cleaved PARP, demonstrated the superiority of **31** (Supporting Information).

To evaluate BBB penetration in vivo, **31** and **32** were given to three mice at 10 mg/kg and 25 mg/kg dosing (analogous to **BP-1-102** in vivo dosing) via intraperitoneal injection, and blood was collected at two time points (30 and 300 min). Brain was also collected from one mouse at each dose and concentrations of **31** and **32** determined by LCMS. We found that after 30 min at 10 mg/kg, compound **31** was found at a concentration of 313 ± 8 nM (Supporting Information). Following these studies, three mice per group were dosed for five consecutive days with 10 mg/kg. Blood was collected at 30 and 300 min post the last dose, and brain was collected from all animals at the 300 min time-point. Then, 313 ± 8 nM of **31**

was detected in the brains of treated animals (Supporting Information). Encouragingly, these studies demonstrated that therapeutic doses of **31** could be achieved in vivo at values similar to the in vitro IC₅₀s demonstrating efficacy against BTSCs.

Next, to evaluate **31** in vivo, NOD-SCID mice were orthotopically xenografted with 10⁵ BT73 glioma cells. On the basis of the results for the above in vivo PK/PD and five-day maximum tolerated dose study, animals were dosed with 10 mg/kg **31**, in order to achieve drug accumulation in the brains of the NOD-SCID mice at doses similar to in vitro IC₅₀s (Supporting Information). Therefore, starting on day 7, mice were given 10 doses of 10 mg/kg intraperitoneal injection of **31** or vehicle control on 4 days on/3 days off schedule. Animals were sacrificed 2 h after the last dose and brain tumors extracted and immunohistochemistry performed for pSTAT3, Ki67 (proliferation), and TUNEL (apoptosis). Analysis of tumors showed decreased tumor cells in **31** treated mice using Hematoxylin/Eosin staining (Figure 5A). Significantly, **31** decreased pSTAT3 expression in tumor cells of treated mice (Figure 5B). Furthermore, **31** appears to decrease proliferation (Figure 5C) and increase apoptosis (Figure 5D) of treated tumors. Thus, in vivo studies strongly suggest in vivo potency and on-target anti-STAT3 activity.

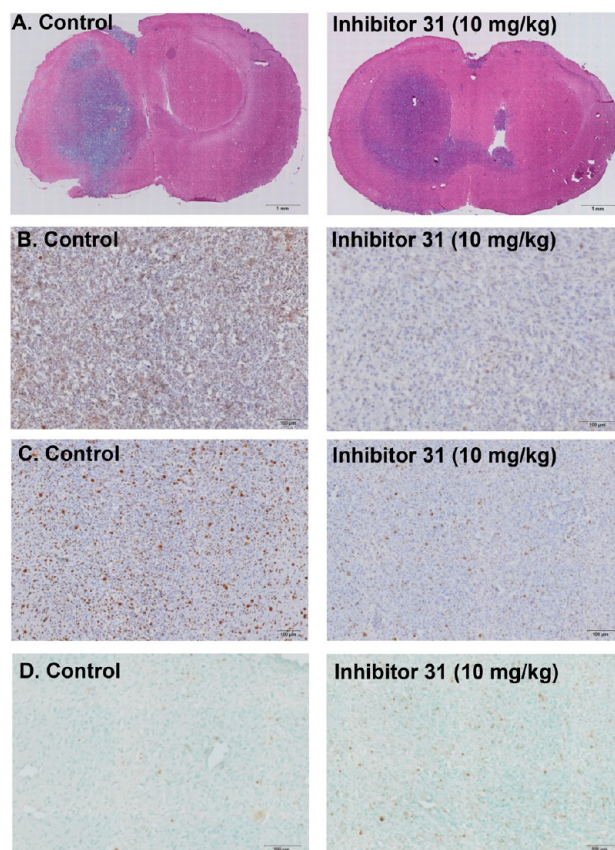


Figure 5. (A) Hematoxylin/Eosin staining: hypercellular tumor dense areas staining blue with hematoxylin. The images display brain tumor suppression and a normal brain morphology for the mice treated with **31**; (B) decreased pSTAT3 in mice orthotopically xenografted with BT73 and treated with **31**; (C) decreased expression of proliferation marker Ki67 in mice treated with **31**; (D) increased apoptosis (TUNEL staining) in mice treated with **31**.

In summary, to the best of our knowledge, we have identified a most potent, nonphosphorylated direct binding STAT3 inhibitor, **31** (SH-4-54). Compound **31** exhibited nanomolar K_D values for STAT3, showed unprecedented cytotoxicity in human BTSCs, displayed no toxicity in human fetal astrocytes, potentially suppressed pSTAT3 with nanomolar IC_{50} s, inhibited STAT3's downstream targets, and showed no discernible off-target effects at therapeutic doses. Moreover, in vivo, **31** exhibits BBB permeability, potently suppresses glioma tumor growth, and inhibits pSTAT3 in vivo. This work, for the first time, demonstrates the power of STAT3 inhibitors for the treatment of BTSCs and validates the therapeutic efficacy of a STAT3 inhibitor for GBM clinical application.

■ ASSOCIATED CONTENT

● Supporting Information

Information on synthesis, characterization, detailed results, and experimental procedures for SPR, Cytotoxicity, BBB permeability, and in vivo assays. This material is available free of charge via the Internet at <http://pubs.acs.org>.

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Author Contributions

*S.H. and H.A.L. contributed equally to this work.

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Notes

The authors declare no competing financial interest.

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