



Published in final edited form as:

*Oncogene*. 2018 March ; 37(11): 1457–1471. doi:10.1038/s41388-017-0049-3.

## LPA signaling is regulated through the primary cilium: a novel target in glioblastoma

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### Abstract

The primary cilium is a ubiquitous organelle presented on most human cells. It is a crucial signaling hub for multiple pathways including growth factor and G-protein coupled receptors. Loss of primary cilia, observed in various cancers, has been shown to affect cell proliferation. Primary cilia formation is drastically decreased in glioblastoma (GBM), however, the role of cilia in normal astrocyte or glioblastoma proliferation has not been explored. Here we report that loss of primary cilia in human astrocytes stimulates growth rate in a lysophosphatidic acid (LPA)-dependent manner. We show that lysophosphatidic acid receptor 1 (LPAR1) is accumulated in primary cilia. LPAR1 signaling through Gα<sub>12</sub>/Gα<sub>q</sub> was previously reported to be responsible for cancer cell proliferation. We found that in ciliated cells, Gα<sub>12</sub> and Gα<sub>q</sub> are excluded from the cilium, creating a barrier against unlimited proliferation, one of the hallmarks of cancer. Upon loss of primary cilia, LPAR1 redistributes to the plasma membrane with a concomitant increase in LPAR1 association with Gα<sub>12</sub> and Gα<sub>q</sub>. Inhibition of LPA signaling with the small molecule compound Ki16425 in deciliated highly proliferative astrocytes or glioblastoma patient-derived cells/xenografts drastically suppresses their growth both *in vitro* and *in vivo*. Moreover, Ki16425 brain delivery via PEG-PLGA nanoparticles inhibited tumor progression in an intracranial

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### Conflicts of interest

The authors declare no conflicts of interest.

**Disclaimers:** This manuscript contains original work only and has not been published or submitted elsewhere. All of the authors have directly participated in the planning, execution and analysis of this study and have approved the submitted version of this manuscript.

glioblastoma PDX model. Overall, our findings establish a novel mechanism by which primary cilium restricts proliferation and indicate that loss of primary cilia is sufficient to increase mitogenic signaling, and is important for the maintenance of a highly proliferative phenotype. Clinical application of LPA inhibitors may prove beneficial to restrict glioblastoma growth and ensure local control of disease.

## Keywords

cilia; proliferation; LPA; GBM; LPAR1

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## Introduction

The primary cilium is a ubiquitous, microtubule-based organelle which is built on top of a membrane-anchored basal body. Primary cilium is an important negative regulator of proliferation and a key sensory organelle. It serves as a hub for multiple signaling cascades including receptor tyrosine kinase (1), Sonic hedgehog (SHH) (2), and G-protein coupled receptors (GPCRs) (3). Disassembly of primary cilium leads to release of the basal body, also called the mother centriole, which is required for mitotic spindle formation and mitosis. Multiple mitotic kinases initiate cilium disassembly including AURKA, Plk1, and Nek2 (4-6). Centrosome sequestration is considered one of the primary mechanisms of negative regulation of proliferation by primary cilium. Nevertheless, cilium shortening and disassembly is often observed immediately after growth factor stimulation in interphase (4). The role of this short-term disassembly in cell proliferation is currently unknown. Recent publications suggest that in different types of cancer including breast, prostate, renal, and glioblastoma (GBM), cilia tend to be lost (7-10). However, the importance of this event on tumor maintenance and progression or treatment is not well understood.

Several studies report that loss of primary cilia in normal cells increases proliferation and supports attachment-independent growth (11, 12), which are common hallmarks of cancer. Attempts to restore primary cilia in cancer cells yield a significant inhibition on proliferation (12, 13). These observations suggest that cilia loss can promote/sustain a highly proliferative phenotype. However, in a subset of Sonic hedgehog (SHH)-dependent medulloblastomas, presence of cilium is mandatory (14) for cancer maintenance, therefore studies on specific cancer subtypes are warranted to establish cilia's role in tumor biology and potential therapeutic applications. Astrocytoma is the most commonly diagnosed adult brain cancer (15), which often progresses to GBM. The majority of GBM patients succumb to the disease within 13-16 months (15). GBM is a highly proliferative disease with limited treatment options (16). Lysophosphatidic acid (LPA) is an abundant mitogen in brain tissue (17). LPA acts through binding of heterotrimeric G-protein coupled receptors (LPAR1-6). It was previously reported that LPAR1 can signal through  $G_{\alpha i}$ ,  $G_{\alpha 12}$ , and  $G_{\alpha q}$  family members (18, 19). LPA stimulates cell proliferation in astrocytes (20) and cancer cells (21). Astrocytes are abundant glial cells and well known for their ability to proliferate, especially in the activated state (22), and were previously reported as the potential cells of origin for GBM (23).

In our current work, we establish that loss of primary cilia promotes proliferation of primary non-transformed human astrocytes, providing permissive conditions for transformation in an LPA-dependent manner. GBM primary cells, with a decreased occurrence of primary cilia, were also sensitive to LPA and LPAR1/3 inhibition. Mechanistically, we found that LPA-LPAR1-driven mitogenic signaling was restricted in cells with primary cilium due to compartmentalization of LPAR1 and its downstream effectors, G $\alpha$ 12 and G $\alpha$ q, in cilia and in cytoplasm, respectively. LPAR1 was redistributed to the plasma membrane upon loss of primary cilium, thus enabling its binding to G $\alpha$ 12 and G $\alpha$ q, and therefore suggesting that redistribution of LPAR1 is a key mechanism driving proliferation in a cilia-dependent manner.

Inhibition of LPAR1/3 with the small molecule inhibitor Ki16425 significantly reduces cell growth rate only in deciliated astrocytes. Likewise, patient-derived GBM proliferation was stimulated by LPA and abrogated by Ki16425 in a dose-dependent manner. Importantly, the growth of GBM patient-derived xenografts *in vivo* was drastically decreased upon Ki16425 administration as a monotherapy without significant side effects.

Taken together, our findings indicate that loss of primary cilia eliminates spatial barriers curbing proliferation, thus unlocking the potential for unlimited proliferation. LPA is one of the key mitogenic factors driving highly proliferative GBM with no or very low basal ciliation, therefore clinical interventions based on inhibiting LPA signaling may significantly improve GBM patient survival and local disease control.

## Results

### Loss of primary cilium promotes proliferation of astrocytes

Human astrocytes (HA) can form primary cilia. Incubation in serum-free media (SFM) promotes ciliation resulting in nearly 80% of cells having cilia (Fig.1A-B). This ciliation rate is similar to that observed *in vivo* (24). To allow for long term experiments, the primary human astrocytes (HA) were immortalized using SV40 large T antigen (HA-LTA) (25). Immortalization by LTA did not affect ciliation, which was similar to the parental astrocytes (Fig.S1A). To test how loss of primary cilia affects proliferation of primary (HA) or immortalized (HA-LTA) astrocytes, we utilized shRNA-driven knockdown of IFT88 or KIF3B, which are well characterized components of cilium assembly machinery (26, 27). Two shRNAs were used to target IFT88 or KIF3B resulting in up to a 90% knockdown (Fig. 1C). Depletion of either IFT88 or KIF3B was sufficient to decrease ciliation to 5-15% (Fig. 1D-E). These ciliation rates are similar to tissue biopsies from GBM patients (28).

The loss of primary cilia resulted in a significant increase in growth rate of both immortalized and primary astrocytes (Fig1.F-G). Interestingly, this difference in growth rate was observed between ciliated (shCon) and deciliated (shKIF3B or shIFT88) cells only upon addition of serum-supplemented media (SSM), but not in SFM. Addition of SSM caused resorption of primary cilia in control cells in a biphasic wave pattern (Fig.S1B), which has been described previously for other cell types (4). These findings indicate that loss of primary cilia by itself does not promote proliferation, but rather increases the response to some mitogenic stimuli present in serum.

### Loss of primary cilia changes spatio-temporal response to mitogen stimulation

To elucidate the potential mechanism underlying cilia-dependent changes in cell growth rate, a time course analysis of synchronized (SFM starved) astrocytes was performed. Phosphorylation of ERK1/2 (Thr202/Tyr204) and AKT (Ser473), common readouts for a variety of mitogen and pro-survival stimuli, was used to follow the serum-induced response. In agreement with previous reports, addition of 10% serum triggers a rapid (5-60min) increase in ERK1/2 and AKT phosphorylation followed by a gradual decrease (60-240min) in phosphorylation in all cell lines independent of ciliation status (Fig.2, Fig.3). Interestingly, the amount of phosphorylated ERK1/2 was twofold higher in deciliated cells (shKIF3B or shIFT88) than in control (shCon) based on immunofluorescent and western blot assays (Fig.2A-F). Contrary to ERK1/2, phosphorylation of AKT was twofold lower in deciliated cells than in control (Fig.3A-F), indicating that changes in the pattern and intensity of the signal initiation/propagation act in a cilia-dependent manner.

### Lysophosphatidic acid signaling is critical for increased proliferation in deciliated astrocytes

To define the mitogene/s potentially responsible for the hyperproliferative phenotype of deciliated astrocytes, charcoal-stripped serum was used. Charcoal stripping is an efficient way to deplete bioactive lipids, as well as hormones and some vitamins (29), but preserve protein-based growth factors. Surprisingly, charcoal-treatment of serum (cSSM) was sufficient to completely abrogate the increase in growth rate observed in deciliated astrocytes (Fig.4A). One of the most abundant lipid-based mitogens found in serum is lysophosphatidic acid (LPA), which binds to LPA receptors (LPARs) to elicit a response. To test if LPA is involved in deciliation-dependent stimulation of proliferation, a small molecule inhibitor of LPAR1-3, Ki16425 (30), was used. Similar to cSSM, addition of Ki16425 to SSM was sufficient to abrogate the deciliation-dependent increase in growth rates, without any effect on growth of ciliated astrocytes (SSM vs. SSM+Ki16425) (Fig.4B). Moreover, addition of LPA alone to the serum-free medium (SFM) was sufficient to stimulate proliferation of astrocytes, and partially recapitulate the difference observed between ciliated and deciliated cells in the presence of serum (Fig.4B). As expected, LPA addition resulted in a pattern of ERK1/2 and AKT phosphorylation similar to the one observed during serum stimulation (Fig.4C-E). Surprisingly, treatment with EGF, bFGF, PDGF-A/B, or HGF as single agents was neither sufficient to induce proliferation (Fig.S2A) nor able to recapitulate the difference in phosphorylation of both ERK1/2 and AKT (Fig.S2B-C, Fig.S3A-B) previously observed between ciliated and deciliated cells (Fig.2E, Fig.3E). Interestingly, addition of a cocktail of protein growth factors (GFs) including EGF, bFGF, and B27 supplement along with LPA was sufficient to fully recapitulate the phenotype observed with addition of serum (Fig.4F). These findings indicate that LPA signaling is responsible for the highly proliferative phenotype observed in astrocytes with disrupted ciliogenesis, but it requires additional growth factors to amplify its effect.

### LPAR1 localizes to primary cilia

To understand the mechanisms underlying cilia-dependent action of LPA, the subcellular localization of LPARs and G $\alpha$  subunits participating in downstream signal transduction

pathways was analyzed. A panel of LPARs fused with 3xFLAG-tag was exogenously expressed in astrocytes followed by immunofluorescent analysis using anti-FLAG antibodies. LPAR1 and LPAR3, but not LPAR6, were consistently localized in primary cilium (Fig.5A, S4A-B). Interestingly, in deciliated cells, both LPAR1 and LPAR3 were targeted to the plasma membrane (Fig.S5A-B). To exclude the possibility of overexpression-driven cilium targeting of LPAR1, we validated several anti-LPAR1 antibodies for immunofluorescent staining, using CRISPR-Cas9 driven LPAR1 depleted astrocytes as a control (Fig.S6). Utilizing validated antibodies we were able to confirm that endogenous LPAR1 localizes to primary cilium when it is present (Fig.S7).

Analysis of the cellular localization of multiple G $\alpha$  subunits shows that only G $\alpha$ s consistently targeted to primary cilia, similar to previous reports (31). However, whether other G $\alpha$  subunits enter primary cilia is currently unknown. We found that G $\alpha$ i1, G $\alpha$ q, and G $\alpha$ 12 did not display ciliary localization, but rather diffuse cytoplasmic and plasma membrane staining (Fig.5C). Such a pattern of compartmentalization suggests that LPARs may engage in interactions with different G $\alpha$  subunits depending upon presence or absence of primary cilia.

#### **LPAR1 directly binds to G $\alpha$ 12 and G $\alpha$ q subunits in deciliated cells**

To test this hypothesis, immunoprecipitation analysis was performed in shCon, shIFT88, and shKIF3B astrocytes transiently overexpressing 3xFLAG-LPAR1. Based on sequence similarity, LPAR1 is predicted to interact with G $\alpha$ i, G $\alpha$ q, and G $\alpha$ 12 (19). No interaction was detected between LPAR1 and G $\alpha$ s. The robust co-immunoprecipitation of G $\alpha$ 12 and G $\alpha$ q with LPAR1 was noted, being 2-3 times higher in deciliated astrocytes (Fig.5D). Previously, LPAR1 was reported to promote cancer cell proliferation specifically through G $\alpha$ 12 (18). G $\alpha$ q was also reported to transmit pro-proliferative signals in cancer (32). We concluded that LPAR1 is sequestered in primary cilia, which prevents its interaction with cytoplasmic G $\alpha$ 12 and G $\alpha$ q, restricting its proliferative signaling. Loss of cilia promotes LPAR1 interaction with G $\alpha$ 12 and G $\alpha$ q, thus promoting the proliferative response to LPA (Fig.5D).

#### **LPA signaling drives GBM proliferation both *in vitro* and *in vivo***

To evaluate our findings in disease-relevant settings, previously characterized GBM patient-derived cells and xenografts (33, 34) of two molecular subtypes (classical-GBM6 and mesenchymal-GBM12) were used. In agreement with previous reports (28), only 5-10% of GBM cells *in vivo* (GBM xenografts in mice, Fig.S8A-B) or *in vitro* (primary cells short term cultures, Fig.S8C-D) possess primary cilia. LPAR1 staining in GBM6 and 12 cells shows similar pattern to primary astrocytes (Fig.S8E). LPA was previously implicated in GBM progression and invasiveness (35), hence its effects on GBM cell proliferation was evaluated. Upon addition of LPA in serum-free GF-supplemented media, a significant increase of primary GBM cell growth rate *in vitro* was observed (Fig.S8F), while addition of the LPA inhibitor, Ki16425, to SSM decreases GBM cell growth rate in a dose-dependent manner (Fig.S8G).

Similar to *in vitro* studies, daily intraperitoneal administration of 30mg/kg Ki16425 as a monotherapy over 4-5 weeks significantly decreased the growth of subcutaneously transplanted GBM xenografts in immunodeficient mice (Fig 6A-E). In agreement with *in vitro* PDX cell line proliferation data (Fig.S8G), the number of mitotic figures in Ki16425-treated tumors (Fig 6D, E) was twofold lower. Overall, these findings indicate that LPA is an important mitogen in GBM and inhibition of LPA signaling is a viable option to improve anti-GBM therapy.

### **PEG-PLGA Ki16425 loaded nanoparticles slow down GBM growth in brain**

To assess the feasibility of targeting GBM with Ki16425 in patients, its effect on GBM growth was evaluated in an intracranial mouse model. Preliminary studies indicated that the regimen used for the subcutaneous GBM PDX experiment was not efficient in the intracranial model (Fig.S9A-B), suggesting that Ki16425 is not capable of crossing the blood brain barrier. To overcome this issue, we used a PEG-PLGA nanoparticle-based delivery system (36) to allow for robust Ki16425 brain entry. Fluorescently labeled nanoparticles rapidly entered the brain upon IP injection and were gradually excreted/ degraded over the next 12h period (Fig.S9C-D). To account for this decay, the regimen was modified to 30mg/kg Ki16425 loaded into nanoparticles and delivered every 12h. Mice were intracranially injected with  $5 \times 10^5$  GBM12 cells and tumor growth was monitored weekly via MRI. Upon tumors reaching 5mm<sup>3</sup>, mice were randomly assigned to Ki16425 or a vehicle loaded nanoparticles control group. Over 2 weeks of treatment, the Ki16425 nanoparticle-treated group showed a twofold decrease in tumor progression, compared to control (Fig.7A-B). In agreement with our previous experiments, the number of mitotic cells was decreased twofold (Fig.7C-D). These findings show that Ki16425 with a proper delivery method can significantly suppress GBM progression, and potentially, in combination with the current standard of care, improve local disease control and GBM patient survival.

## **Discussion**

Primary cilium is well known for its role in multiple signaling cascades (1, 2, 11). Primary cilia length and the number of cells with cilium is significantly reduced or lost in multiple cancer types including GBM (7-10). Moreover, several studies have noted an increase in cell proliferation upon loss of primary cilia (11, 12). The proposed mechanism of cilia-driven proliferation control involves the sequestration of basal body (mother centriole) and the inability to form the mitotic spindle (37). Previously, it was shown that primary cilium disassembly is biphasic: a first wave of fast, transient disassembly within 1-2h after exposure to mitogenic stimuli (4), followed by a second wave of disassembly at 18-24h, which coincides with mitotic entry. The role of the first wave of cilium disassembly in mitogenic signaling is currently unknown, as is the mechanism/s underlying the increase in proliferation upon loss of primary cilia.

Several pro-proliferative signaling cascades are reported to require primary cilium for proper signal transduction. For instance, PDGF-AA signaling through PDGFR $\alpha$  is lost in deciliated fibroblasts, therefore stimulation with PDGF-AA does not cause ERK1/2 or AKT activation (38); however, the overall proliferation effect of cilia loss was not evaluated in this

work. SHH signaling is well-studied in conjunction with primary cilium. Loss of cilia ablates SHH signaling and SHH-driven proliferation (39), and correspondingly, SHH-driven cancers have a tendency to maintain high ciliation rates (14, 40). On the other hand, ciliary localization of IGF1R (41) and Notch (11) signaling is needed for differentiation and restriction of cell proliferation. Importantly, cilium can selectively sequester signaling components like GPR88 (42) and LPAR1/3 (Fig.5, S4) from their interaction with cilia-excluded co-factors, thus conferring selectivity on signal propagation/amplification through the lateral segregation of receptors. This strongly argues that the effect of primary cilia on proliferation is context-dependent, with pro-proliferative effects during development through SHH signaling and anti-proliferative effects in more differentiated cells.

In GBM, the complete inhibition of ciliogenesis seems to have minimal and non-coherent effects on overall proliferation and tumor progression (43). Nevertheless, ciliation compared to normal brain cells is drastically decreased (9), supporting an anti-mitogenic function of primary cilium in GBM. Our current findings suggest that the engineered loss of primary cilia in normal human astrocytes phenotypically closely mimics transient cilium disassembly (Fig.S1B), and also resembles cilia loss in GBM cells (Fig.S8B, D). The increase in growth rate observed in deciliated cells (Fig.1F, G) suggests that transient cilia disassembly may be a key event augmenting mitogenic signaling. Mechanistically, we found that loss of primary cilia in human non-transformed astrocytes results in the amplification of ERK1/2 phosphorylation and promotes proliferation in an LPA-dependent manner (Fig.2, Fig.4). Interestingly, phosphorylation of AKT in response to LPA or serum stimulation was decreased in deciliated astrocytes (Fig.3). This can be attributed to inactivation of PDGFR $\alpha$  (38) and IGF1R (41) driven signaling, and adds to the understanding of increased stress sensitivity in deciliated astrocytes (44). These findings support the idea that loss of primary cilia changes the pattern of the cellular response to mitogen stimulation, resulting in higher, more sustainable ERK1/2 activation thus explaining the increase in proliferation, but revealing the potential vulnerability of deciliated cells to stress via a decrease in pro-survival signaling (pAKT). Since GBMs usually have high pAKT levels (45) and we did not observe the transformation of deciliated astrocytes, we conclude that additional hit/s such as those well-known for GBM including RTK amplification/PTEN inactivation/PI3K activation, are required for overcoming the decrease in AKT activation.

Previously, LPA was implicated in the regulation of cell migration and proliferation (21, 35), however, the role of primary cilia in these signaling pathways was not explored. The LPA receptors (LPAR1/3) are specifically localized to primary cilium, but the LPAR downstream effectors, G $\alpha$ 12 and G $\alpha$ q, are excluded from it (Fig.5, S4). Both G $\alpha$ 12 and G $\alpha$ q activation is known to be implicated in cancer cell proliferation and cancer progression (46, 47). For G $\alpha$ 12, activation through LPAR1 specifically was reported in ovarian cancer cells (18). Our findings support the notion that in the absence of primary cilia, LPAR1 localizes to the plasma membrane and interacts with G $\alpha$ 12 and G $\alpha$ q (Fig.5D, S5, S7). However, further research is required to pinpoint the intracellular compartment for G $\alpha$ 12 and G $\alpha$ q and endogenous LPAR1 interaction. Hence, loss of primary cilia could be responsible for increased proliferation in a subset of cancers, and thus cilia restoration or the manipulation of cilia-dependent signaling such as LPA could be used to develop new therapeutic approaches to fight cancer.

GBM cells seem to be highly dependent on LPA as a mitogen, since inhibition of LPA signaling with Ki16425 abrogates their growth *in vitro*, and in both subcutaneous and intracranial GBM models (Fig.S8, Fig.6, Fig.7). Interestingly, the magnitude of the Ki16425 effect was the same between two *in vivo* models (Fig.6, Fig.7), suggesting that LPA signaling is equally engaged in both of them. LPA is known to be highly abundant in brain (3.7-35 pmol/mg) and serum concentration of LPA was reported to be close to this range (4-15.5  $\mu$ M) (17). In addition, GBMs were reported to increase LPA production by secreting autotaxin (48). However, further studies are required for comprising the LPA levels in GBM PDXs grown in subcutaneous versus brain settings.

GBM6 and GBM12 used in the current study are of classical and mesenchymal molecular subtypes respectively, and are highly aggressive in mouse models (33, 34). Both of these subtypes are associated with rapid disease progression and poor prognosis. Our *in vivo* experiments clearly show that targeting LPA signaling yields a twofold reduction of tumor growth, specifically through a decrease in the proliferation rate of tumor cells (Fig.6, Fig.7). Moreover, utilizing a PEG-PLGA nanoparticles delivery system confirmed the effectiveness of targeting LPA signaling in intracranial settings. The median survival of GBM patients with the current standard of care including aggressive surgery, radiation, and chemotherapy (49) is about 12 months (15), with the majority of patients experiencing recurrence within 32-36 weeks (16). Targeting LPA signaling can prove to be highly beneficial in addition to standard care, since LPA is implicated in proliferation and migration/invasion (21, 35). Further studies are required, but based on our findings, we expect a substantial increase in recurrence-free survival upon inhibition of LPA signaling.

Overall, our data supports the role of LPA signaling in cancer cell proliferation, and for the first time, highlights primary cilia as a switch for the interpretation of LPA as a mitogen.

## Materials and Methods

### Cell lines and reagents

Human astrocytes isolated from human cortex (1800) were obtained from ScienCell Research Labs and maintained in DMEM/F12, supplemented with 10% heat-inactivated FBS, Antibiotic-Antimycotic (ThermoFisher) and 10  $\mu$ g/ml of gentamycin (MP Biomedicals). Cells were propagated and cryopreserved at passage 2, for all studies cells were not passaged more than 10 times or 8 weeks, no authentication or mycoplasma testing were performed. For immortalization, primary astrocytes at passage 5 or 6 were cultured until 50–75% confluency and transfected with SV40 large T-antigen construct (25), followed by selection with 500 $\mu$ g/ml G418 (Sigma) until resistant colonies were formed. GBM6 and 12 PDXs (33) were kindly provided by Dr. Jann Sarkaria (Mayo Clinic, Rochester, MN) through a shared MTA agreement, no authentication or mycoplasma testing were performed. PDXs were maintained and used for intracranial injections or primary cell culture isolation as previously described (34). A list of antibodies and their used applications is outlined in Table S1. shRNA constructs were obtained from Dharmacon and are outlined in Table S2. All primers were purchased from IDT Technologies or Invitrogen, the sequences of which are reported in Table S3. The following reagents were used: Lysophosphatidic acid (LPA, Cayman Chemical), EGF and PDGF-AB (Sigma), HGF (R&D Technologies), bFGF



(Peprotech), B27 supplement (ThermoFisher), Ki16425 (ApexBio). For lysophosphatidic acid stimulation experiments, bovine serum albumin (BSA; BP1600, Fisher Scientific) was used as a carrier and was added to all cells to a final concentration of 0.1%.

### Western blotting

Western blotting was performed using standard procedures (50). Primary antibodies used are outlined in Table S1. Secondary anti-mouse and anti-rabbit HRP-conjugated antibodies (Jackson ImmunoResearch Labs) were diluted 1:10,000 followed by chemiluminescence-based detection with HyGLO™ (Denville Scientific). Bands were quantified using the digital electrophoresis documentation and image analysis software GeneTools (Syngene Corp.), with signal intensity normalized to either  $\alpha$ -tubulin or GAPDH.

### Immunofluorescent cell analysis

Cells were processed as previously described (4). Primary antibodies used are outlined in Table S1. Secondary antibodies included AlexaFluor 488, 555, and 647 (ThermoFisher). Images were captured using a standard setting by an LSM510 confocal microscope (Zeiss) (50). All images represent whole-cell 3D reconstructed projections with 0.4 $\mu$ m steps. All quantifications were done using ImageJ (NIH).

### Cell growth/proliferation analysis

Cell growth rate was determined using semi-automated cell counting in ImageJ (NIH). Cells were plated at  $1 \times 10^4$  cells per well and grown in the specified conditions for five days, fixed with methanol, and stained with Hoechst33342. Four random 10x fields per well were analyzed with at least three replicates per independent experiment; each graph represents at least three independent experiments.

### Fluorescent immunohistochemistry (F-IHC)

Deparaffinization and rehydration of 4-5 $\mu$ m thin sections was performed as following: 1) three times for 3 min in xylene, 2) three times for 2 min in 100% ethanol, 3) 2 min in 95% ethanol, 4) 2 min in 80% ethanol, 5) 2 min in 70% ethanol, and 6) 5 min in 1XTBS. Antigen retrieval was done using 98°C citrate buffer, pH 6.0 for 20 minutes. Sections were blocked for 60 min with 5% BSA, 1XTBS solution and stained with the indicated primary antibodies. Secondary antibodies included AlexaFluor-488, 555, and 647 (ThermoFisher) and sections were mounted with ProLong Gold DAPI-containing media (ThermoFisher). Images were captured using LSM510 confocal microscope as previously described (50) (Zeiss). All images represent whole-cell 3D reconstructed projections with 0.4 $\mu$ m steps. All quantifications were done using ImageJ (NIH).

### LPAR cloning

For robust expression and detection of LPARs, pcDNA3.1 vector was modified as previously described (51). Briefly, the insert containing the cleavable ER-targeting sequence followed by a 3xFLAG-tag and multiple cloning site (Table S3) was synthesized through String™ service (ThermoFisher) and cloned into pcDNA3.1 between HindIII and Sall restriction sites. LPAR1, 3, 6 cDNA was amplified from a Human Mammary Epithelial Cells (HMECs)

cDNA library kindly provided by Dr. Alexey Ivanov (West Virginia University, Morgantown, WV); primers are shown in Table S3. The generated inserts were cloned into pcDNA3.1 ER-3xFLAG between BamHI and XhoI restriction sites. All constructs were validated by sequencing.

### Immunoprecipitation

For LPAR1 immunoprecipitation, 3xFLAG-LPAR1 was transiently overexpressed in cells pre-incubated for 24h in serum-free media. Cells were lysed as previously described (52). 3xFLAG-LPAR1 was precipitated with anti-FLAG M2 affinity gel (Sigma) and used for subsequent Western blot analysis.

### Animal experiments

NOD.Cg-*Prkdc<sup>scid</sup> Il2rg<sup>tm1Wjl</sup>/SzJ* (NSG) immunodeficient male mice (The Jackson Laboratory) were housed in the West Virginia University Animal Facility under pathogen-free conditions with an approved Institutional Animal Care and Use Committee protocol. For subcutaneous injections, 100µl of GBM6 or GBM12 tumor mash mixed 1:1 with Matrigel (Corning) was injected into the mice flanks. For intracranial injections, transient primary cell cultures were established as previously described (34), and  $5 \times 10^5$  cells in 5µl of Ca/Mg-free Dulbecco modified Phosphate Buffer Saline (DPBS) were administered into the mouse cortex via stereotactic device-guided injection (34). Subcutaneous tumor growth was assessed weekly via caliper measurements; treatment was initiated upon tumor volume reaching 100mm<sup>3</sup>. Intracranial tumor volume was assessed weekly via contrast enhanced MRI (53) or twice a week via bioluminescence imaging (54); treatment was initiated upon tumor volume exceeding 5mm<sup>3</sup>. Animals were randomly assigned to the treatment group by simple randomization, investigator was single blinded during group allocation. Mice bearing subcutaneous tumors were intraperitoneally injected with 30mg/kg Ki16425 in 95% corn oil/5% DMSO or vehicle alone daily. For the intracranial model, mice were administered 30mg/kg Ki16425 loaded into PEG-PLGA nano-particles in PBS or given nano-particles alone twice daily. All animals were euthanized upon reaching a moribund condition, according to the WVU IACUC Tumor Development and Tumor Scoring in Rodents policy.

### Contrast enhanced Magnetic Resonance Imaging (MRI)

To visualize the intracranial tumors, mice were intraperitoneally injected with gadolinium-DTPA (contrast reagent, BioPAL) to a final concentration of 2mmol/kg (55), and imaged with a compact MRI system (ICON, Bruker). Images were taken with T1 weighted RARE sequence (echo time: 26.23ms; repetition time 1984.158ms; averages: 8; rare factor: 6). Overall time of scan was 25min and final resolution was 0.125mm for all axes. Stack images were reconstructed and analyzed using ImageJ (NIH).

### Animal bioluminescence imaging

Mice were imaged twice a week for quantitative evaluation of tumor growth as previously described (56). Images were obtained using the IVIS Lumina-II Imaging System and Living Image-4.0 software (PerkinElmer).

## PEG-PLGA nano-particles

Nano-particles were prepared as previously described (36). Briefly, PEG-PLGA and Ki16425 were dissolved in acetone and added dropwise into water. The nano-particles were stirred for 2 h at 40°C before being collected by centrifugation (4000g for 90 min). After discarding the supernatant, the nanoparticles were resuspended in Phosphate-Buffered Saline (PBS).

## Statistical analysis

Statistical comparisons were made using two-tailed Student's *t*-test. When more than two groups were analyzed, one-way or two-way analysis of variance (ANOVA) was used.  $P < 0.05$  was considered to be significant (\*) as indicated in figure legends. All treatment groups were compared to vehicle/control unless mentioned otherwise. Experimental values were reported as the means with +/-S.E.M (standard error of mean). All calculations of statistical significance were made using GraphPad software. Sample sizes and statistical analysis chosen were based on our previous experience and recommendations of biostatistician (4, 50, 56).

## Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

## Acknowledgments

The authors thank Dr. Alexey Ivanov for sharing of the HMLE cDNA library and Brandon Jones for invaluable help with manuscript preparation.

**Financial support:** This research was supported by grants R21CA208875 (E.N.P), R01CA148671 (E.N.P) from the NIH/NCI, U54GM104942 (W.J.G) from the NIGMS/NIH, WVU Cancer Institute Undergraduate Research Fellowship (C.L.G), and "Let the Journey Begin" fund established by Erin Dunmire, WVU Cancer Institute. Andrey Bobko is supported by startup funding from WVCTSI. Naira V. Margaryan is supported by U54GM104942 from WVCTSI.

West Virginia University Microscope Imaging Facility is supported by the WVU Cancer Institute and NIH grants P20RR016440, P20 RR016477 and P30RR032138/P30GM103488.

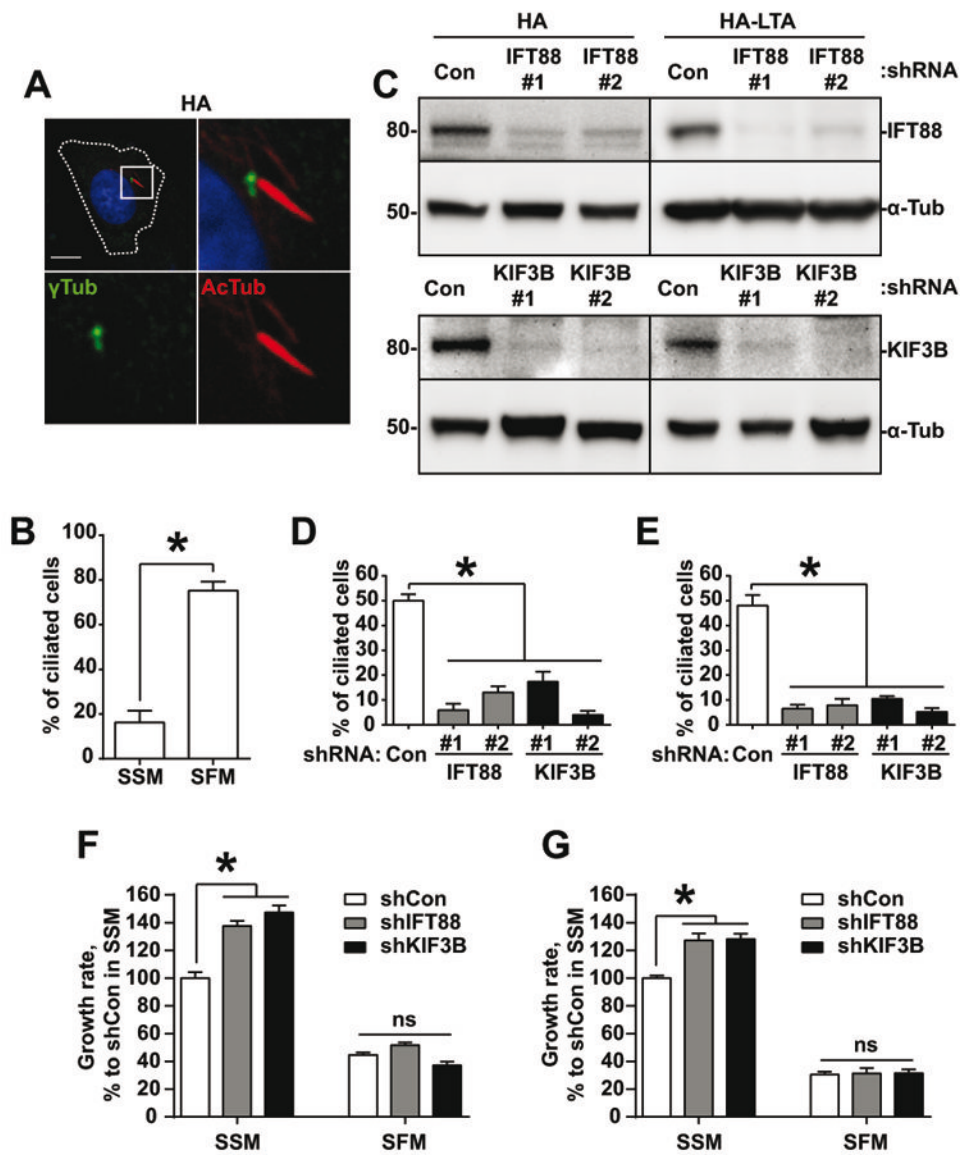
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**Fig. 1. Loss of primary cilia promotes astrocyte proliferation in a growth factor-dependent manner**

(A) Representative image of the cilium formed by primary human astrocyte (HA), stained for acetylated  $\alpha$ -tubulin (AcTub, cilium marker) and  $\gamma$ -tubulin ( $\gamma$ Tub, basal body marker); scale bar – 10 $\mu$ m. (B) Quantification of primary astrocytes forming cilium, as in (A), in regular serum-supplemented media (SSM) or upon 48h of serum starvation (SFM); 300 cells, 100 cells in each of 3 independent experiments; Student's *t*-test,  $p < 0.05$ . (C) Western blot of IFT88 and KIF3B in primary and immortalized astrocytes (HA-LTA) stably expressing non-targeting shRNA (Con) or shRNA against IFT88 or KIF3B. (D, E) Quantification of primary astrocytes (D) and immortalized astrocytes (E) forming primary cilium, as in (B) upon depletion of IFT88 or KIF3B, as in (C); 300 cells, 100 cells in each of 3 independent experiments; one-way ANOVA with Dunnett's post hoc test,  $p < 0.05$ . (F, G) Growth rates of primary astrocytes (F) and immortalized astrocytes (G) upon depletion of

IFT88 or KIF3B in full media or in serum-free conditions (SFM); 3 independent experiments; one-way ANOVA with Dunnett's post hoc test,  $p < 0.05$ .

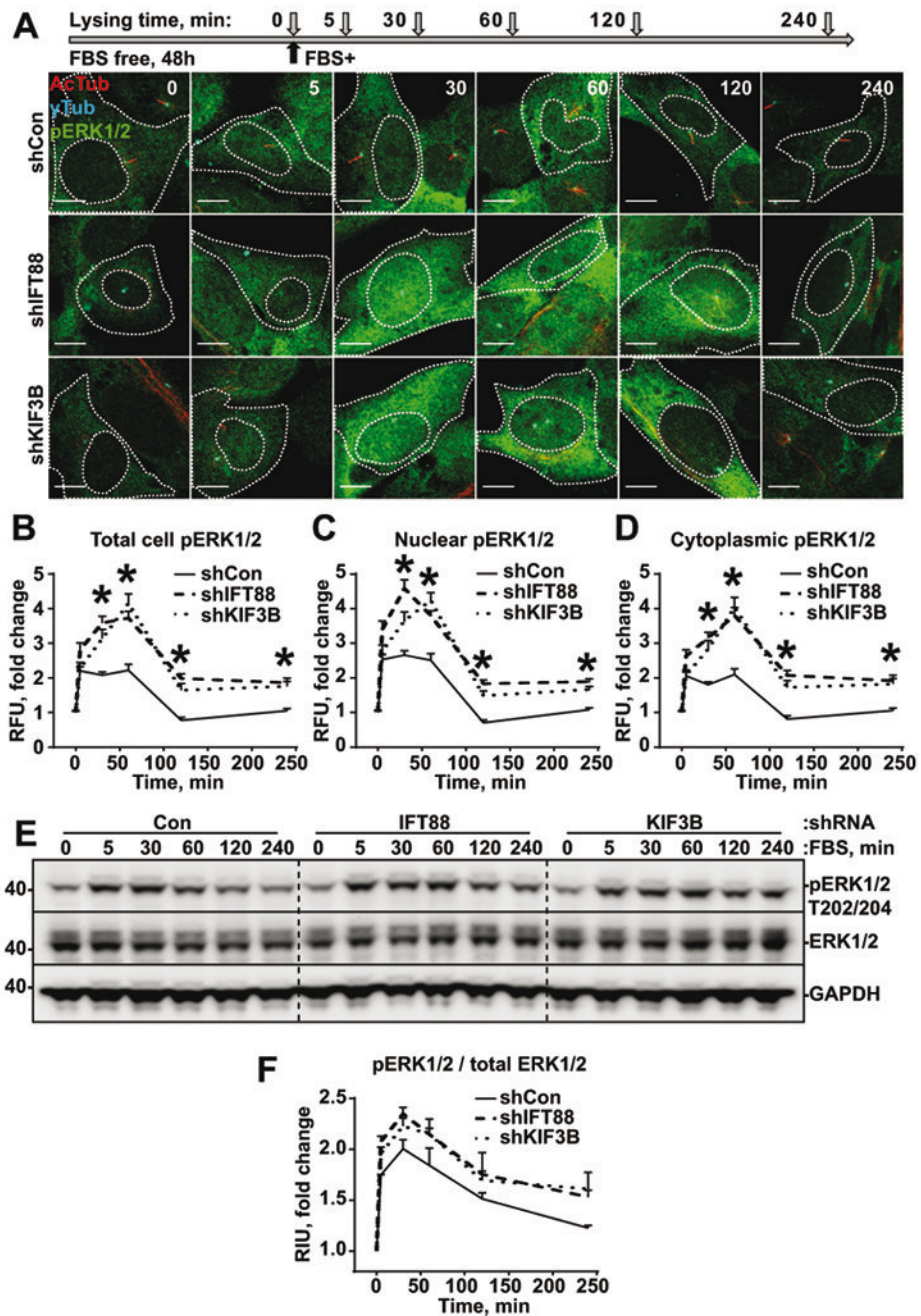
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**Fig. 2. Loss of cilium promotes increased ERK1/2 phosphorylation in response to serum stimulation**

(A) Experiment schematics (top) and representative images of cells stably expressing non-targeting shRNA (Con) or shRNA against IFT88 or KIF3B, stained for acetylated  $\alpha$ -tubulin (AcTub, cilium marker),  $\gamma$ -tubulin ( $\gamma$ Tub, basal body marker), and ERK1/2 phosphor-T202/Y204 (pERK1/2); scale bar – 10 $\mu$ m. (B, C, D) Quantifications of ERK1/2 phosphor-T202/Y204 intensities in whole cells (B), nuclei (C) and cytoplasm (D) as in (A); 100 cells in 3 independent experiments; two-way ANOVA with Dunnett’s post hoc test,  $p < 0.05$ . (E) Representative western blot analysis of cells as in (A), stained for ERK1/2 phosphor-T202/

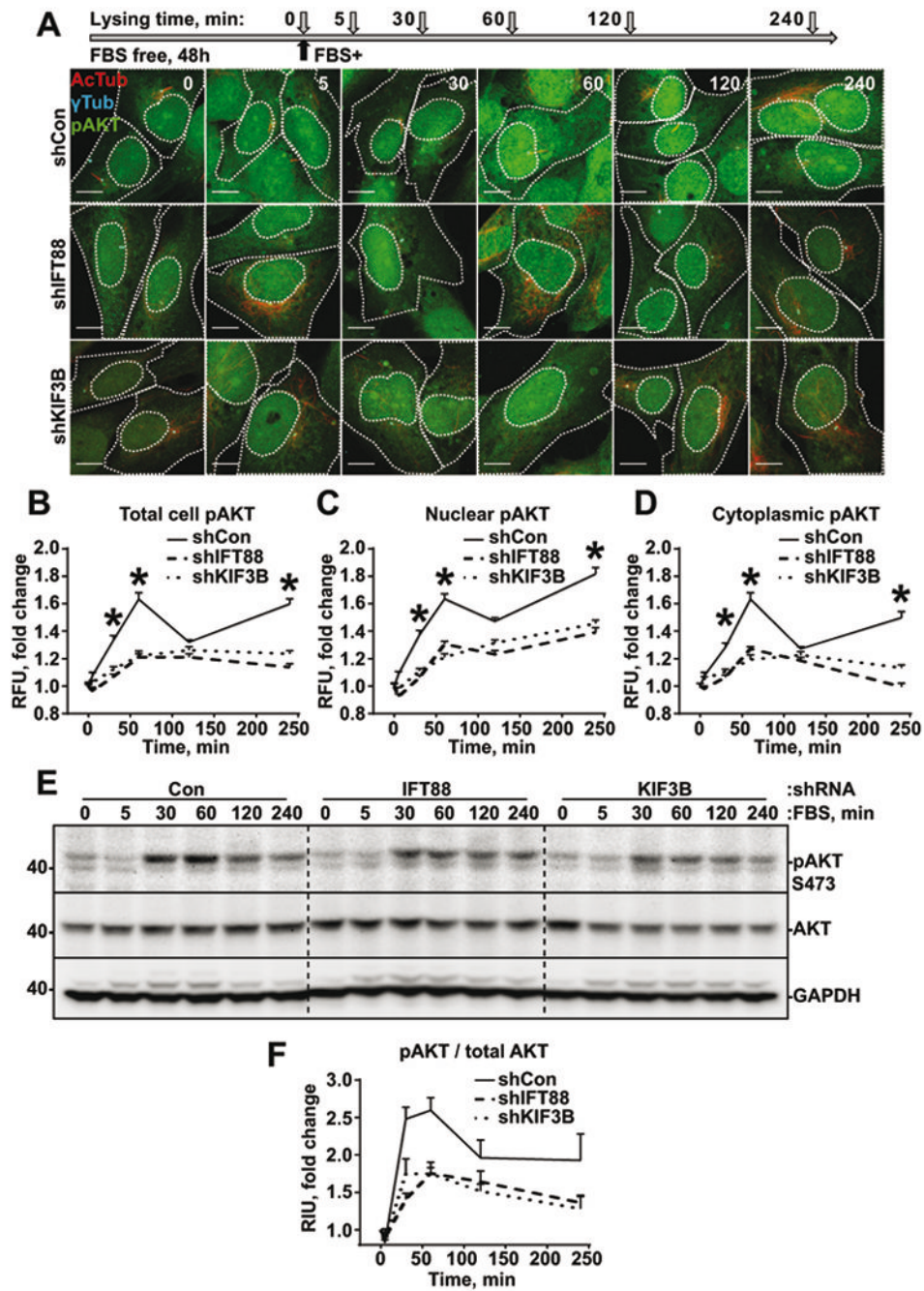
Y204, total ERK1/2, and GAPDH. (F) Quantifications of ERK1/2 phosphor-T202/Y204 bands intensities, as in (E).

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**Fig. 3. Loss of cilium promotes decreased AKT phosphorylation in response to serum stimulation** (A) Experiment schematics (top) and representative images of cells stably expressing non-targeting shRNA (Con) or shRNA against IFT88 or KIF3B, stained for acetylated  $\alpha$ -tubulin (AcTub),  $\gamma$ -tubulin ( $\gamma$ Tub), and AKT phosphor-S473 (pAKT); scale bar – 10 $\mu$ m. (B, C, D) Quantifications of AKT phosphor-S473 intensities in whole cells (B), nuclei (C) and cytoplasm (D) as in (A); 100 cells in 3 independent experiments; two-way ANOVA with Dunnett’s post hoc test,  $p < 0.05$ . (E) Representative western blot analysis of cells as in (A),

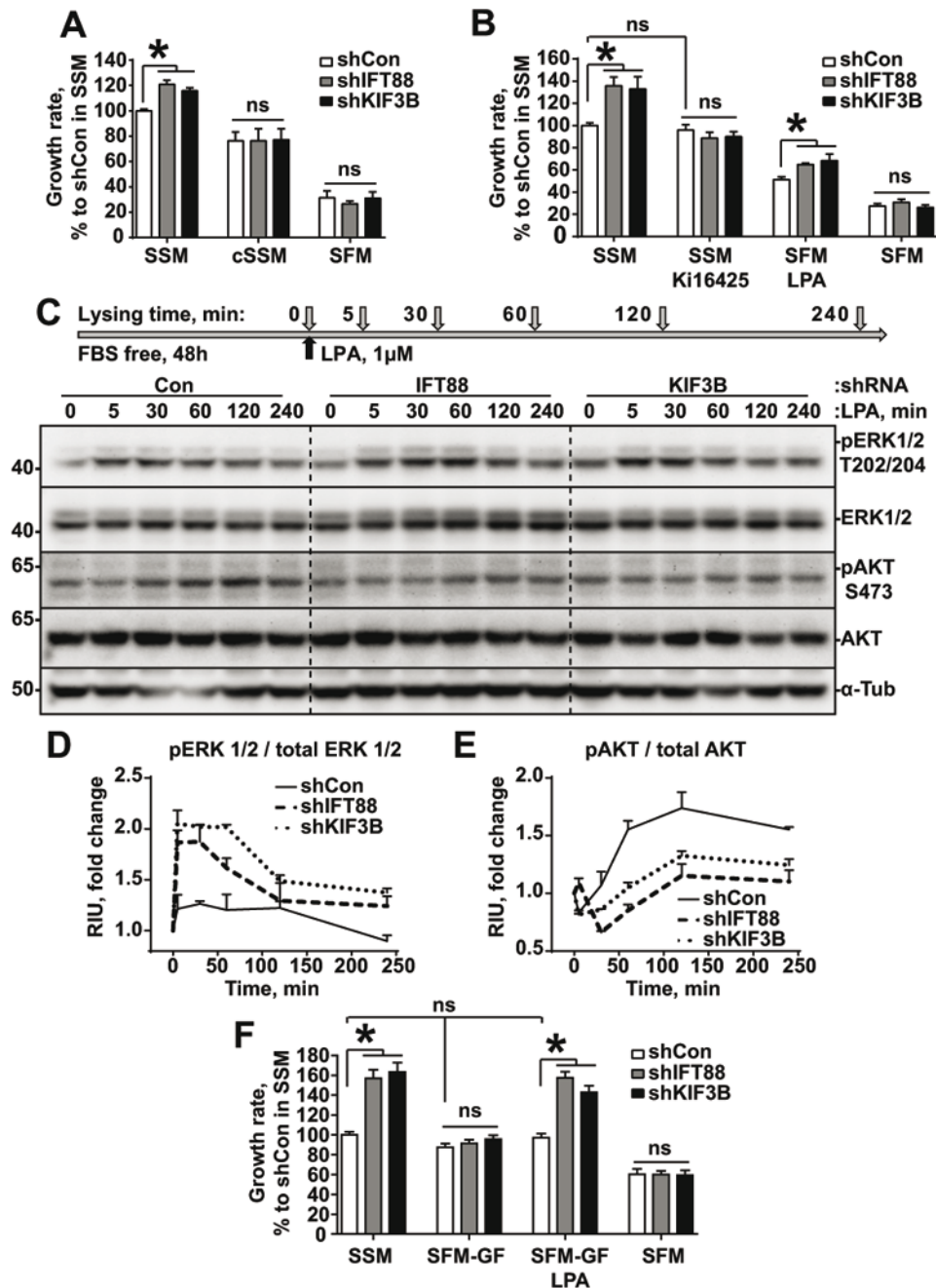
stained for AKT phosphor-S473, total AKT, and GAPDH. (F) Quantifications of AKT phosphor-S473 bands intensities, as in (E).

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**Fig. 4. Lysophosphatidic acid promotes proliferation in deciliated astrocytes**

(A) Growth rates of immortalized astrocytes stably expressing non-targeting shRNA (Con) or shRNA against IFT88 or KIF3B in serum-supplemented media (SSM), media supplemented with charcoal-stripped serum (cSSM), or in serum-free conditions (SFM); 3 independent experiments; one-way ANOVA with Dunnett's post hoc test,  $p < 0.05$ . (B) Growth rates of cells as in (A) in serum-supplemented media (SSM), serum-supplemented media supplemented with 10 μmol/L Ki16425, serum-free media supplemented with 1 μmol/L LPA (SFM LPA), or in serum-free conditions (SFM); 3 independent experiments; one-way ANOVA with Dunnett's post hoc test,  $p < 0.05$ . (C) Experiment schematics (top) and

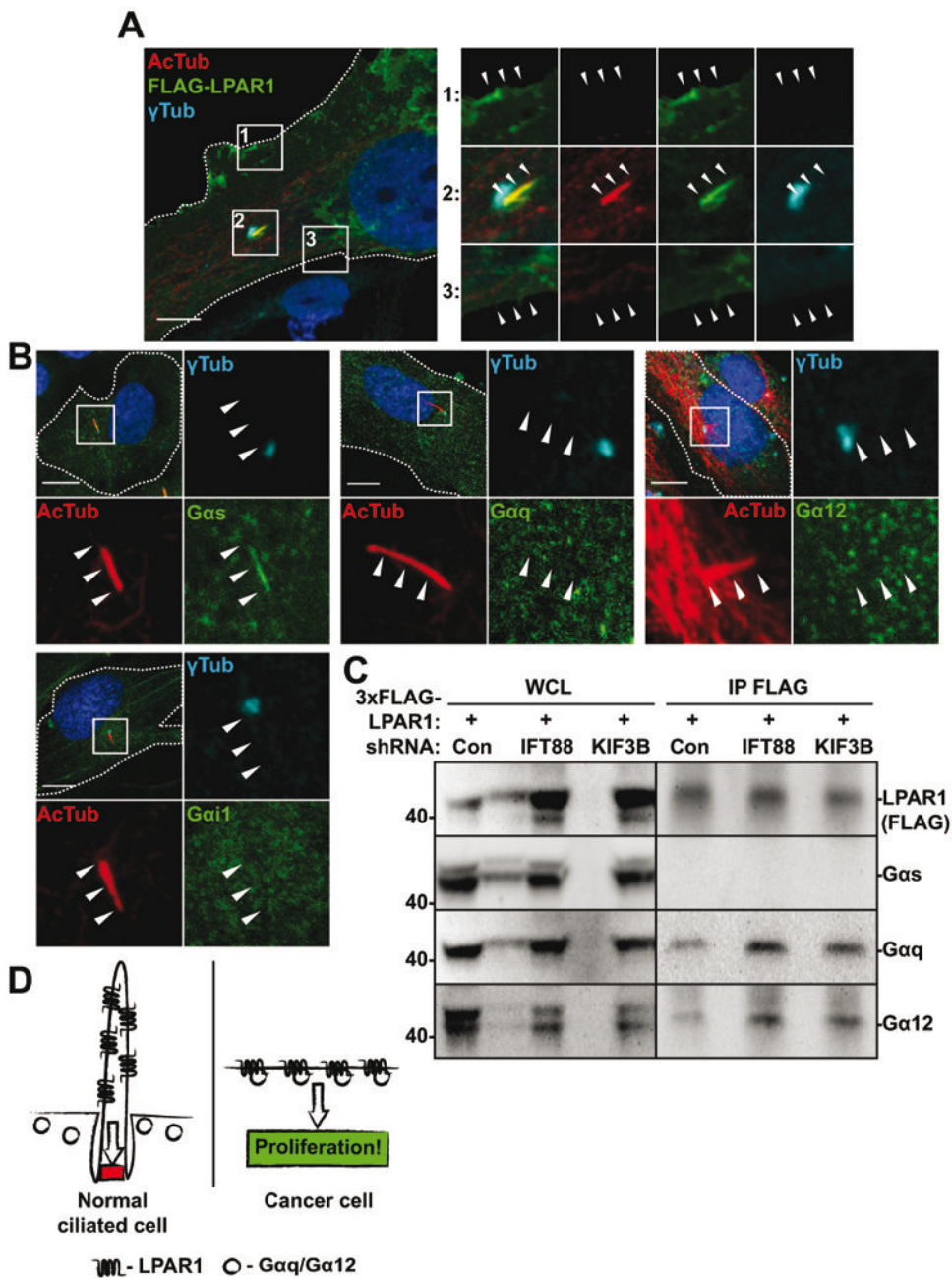
representative western blot analysis of cells as in (A) stained for ERK1/2 phosphor-T202/Y204, total ERK1/2, AKT phosphor-S473, total AKT, and  $\alpha$ -tubulin. (D) Quantifications of ERK1/2 phosphor-T202/Y204 bands intensities, as in (C). (E) Quantifications of AKT phosphor-S473 bands intensities, as in (C). (F) Growth rates of cells as in (A) SSM, SFM supplemented with 20ng/ml EGF, 20ng/ml bFGF, and B27 supplement (SFM-GF), SFM supplemented with 20ng/ml EGF, 20ng/ml bFGF, B27 supplement and 1 $\mu$ mol/L LPA (SFM-GF LPA), or in serum-free conditions (SFM); 3 independent experiments; one-way ANOVA with Dunnett's post hoc test,  $p < 0.05$ .

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**Fig. 5. Intracellular localization of LPA signaling cascade components**

(A) Representative images of immortalized astrocytes expressing exogenous 3xFLAG-LPAR1, stained for acetylated  $\alpha$ -tubulin (AcTub, cilium marker),  $\gamma$ -tubulin ( $\gamma$ Tub, basal body marker) and FLAG-tag, arrowheads indicate primary cilium; scale bar – 10 $\mu$ m (B) Representative images of immortalized astrocytes stained for acetylated  $\alpha$ -tubulin (AcTub),  $\gamma$ -tubulin ( $\gamma$ Tub), Gas, Gaq, Ga12 and Gai1, arrowheads indicate primary cilium; scale bar – 10 $\mu$ m. (C) Immunoprecipitation of 3xFLAG tagged LPAR1 expressed in immortalized astrocytes stably expressing non-targeting shRNA (Con) or shRNA against IFT88 or KIF3B

in serum-free conditions. (D) Schematic of potential mechanism of primary cilium restrictive action on the proliferative component of LPA signaling.

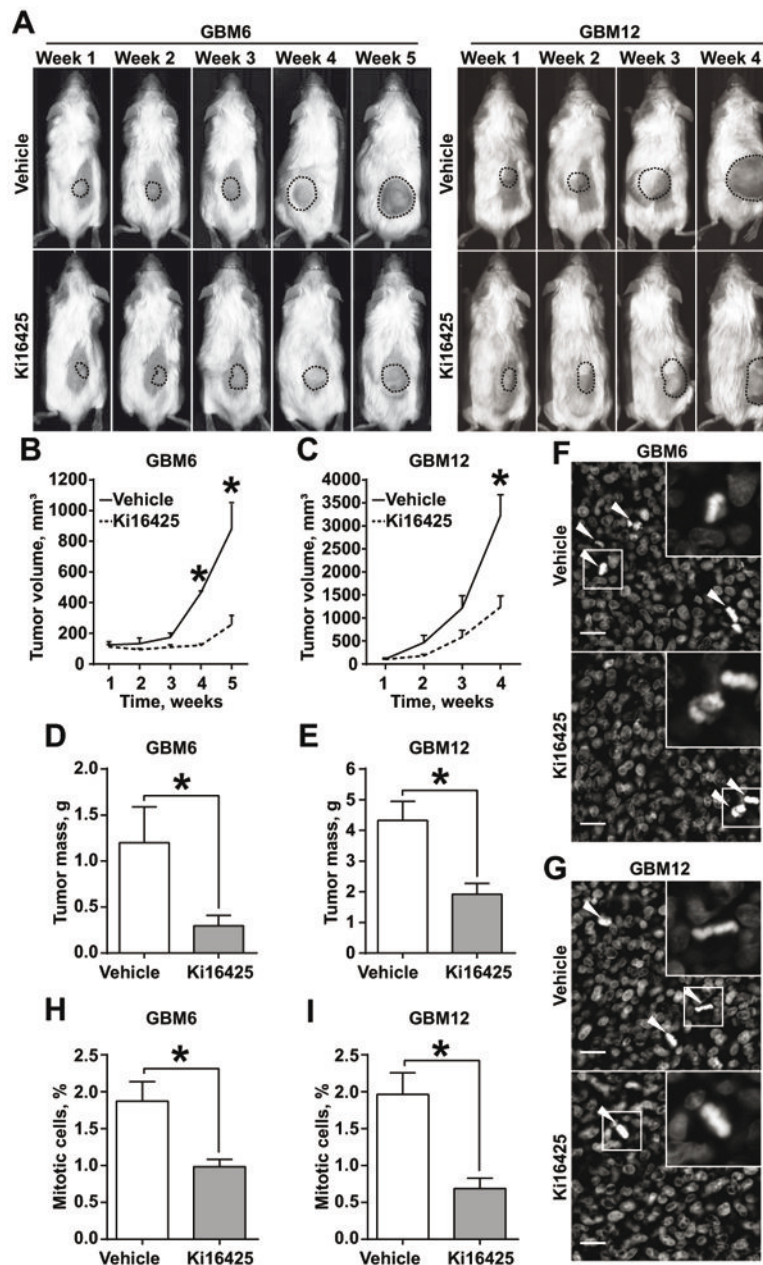
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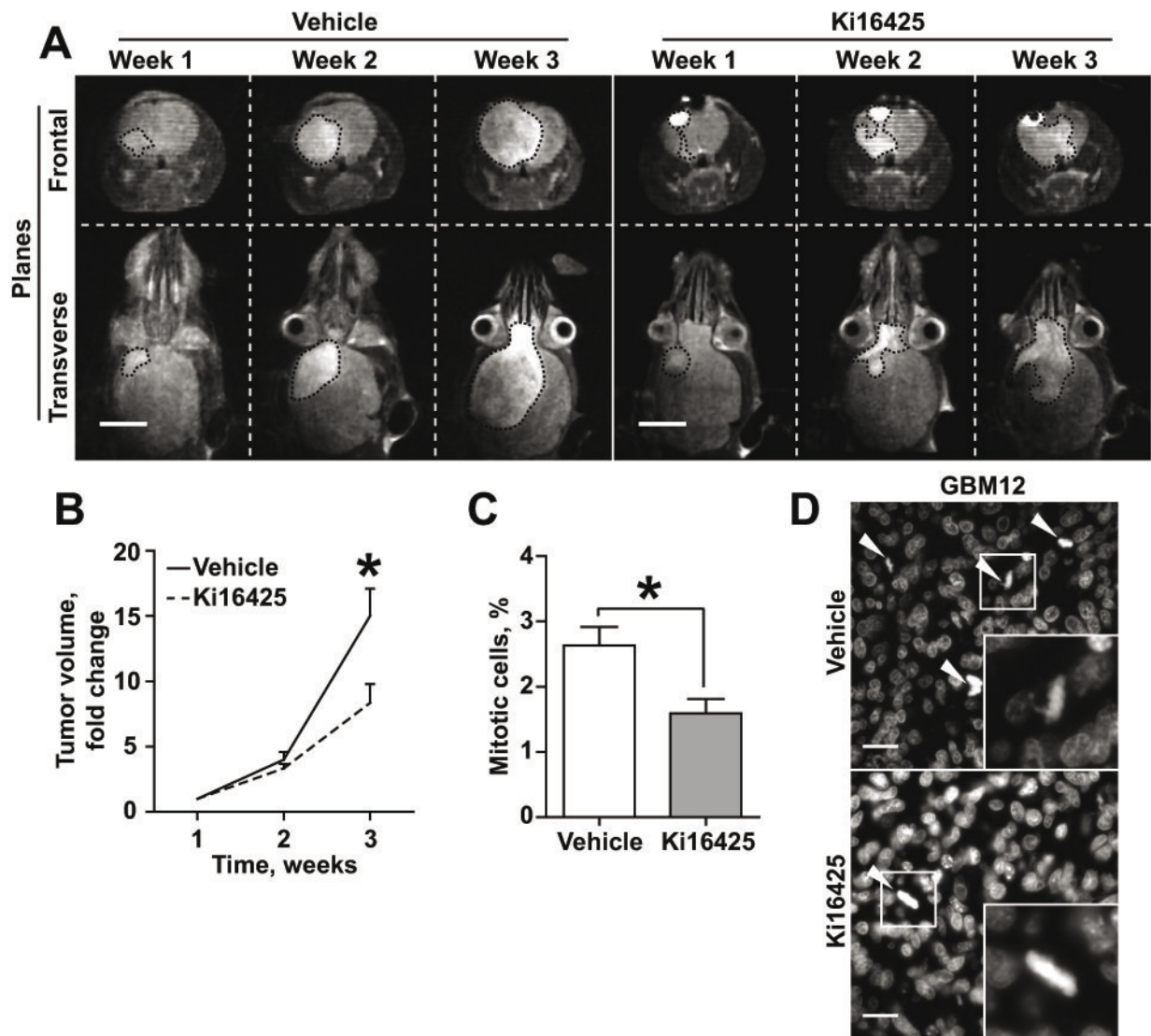
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**Fig. 6. Inhibition of LPA signaling suppresses proliferation of GBM PDXs *in vivo***  
 (A) Representative images of mice subcutaneously injected with GBM6 (left panel) and GBM12 (right panel) and administered with vehicle or 30mg/kg/day Ki16425. (B, C) Analysis of tumor growth as in (A) for GBM6 (B) and GBM12 (C); 5 mice per group; two-way ANOVA with Sidak's post hoc test,  $p < 0.05$ . (D, E) Analysis of GBM6 (D) and GBM12 (E) terminal tumor weight; 5 tumors per group; Student's  $t$ -test,  $p < 0.05$ . (F, G) Representative images of GBM6 (F) and GBM12 (G), stained with DAPI; arrowheads indicate mitotic cells; scale bar – 20 $\mu$ m. (H, I) Quantification of mitotic figures as in (F, G) for GBM6 (H) and GBM12 (I); at least 1000 cells within 10 random fields per group; Student's  $t$ -test,  $p < 0.05$ .



**Fig. 7. Targeted brain delivery of Ki16425 suppresses GBM PDX growth in an intracranial model**

(A) Representative MRI images of mice bearing intracranially grafted GBM12 throughout the treatment with PEG-PLGA nanoparticles loaded with Ki16425 (30mg/kg twice a day); scale bar – 5mm. (B) Analysis of tumor growth as in (A); 5 mice per group; two-way ANOVA with Sidak's post hoc test,  $p < 0.05$ . (C) Quantification of mitotic figures as in (D); at least 1000 cells within 10 random fields per group; Student's  $t$ -test,  $p < 0.05$ . (D) Representative images of tumors from (A), stained with DAPI; arrowheads indicate mitotic cells; scale bar – 20 $\mu$ m.