

AKT1 Activation is Obligatory for Spontaneous BCC Tumor Growth in a Murine Model that Mimics Some Features of Basal Cell Nevus Syndrome

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Abstract

Patients with basal cell nevus syndrome (BCNS), also known as Gorlin syndrome, develop numerous basal cell carcinomas (BCC) due to germline mutations in the tumor suppressor *PTCH1* and aberrant activation of Hedgehog (Hh) signaling. Therapies targeted at components of the Hh pathway, including the smoothened (SMO) inhibitor vismodegib, can ablate these tumors clinically, but tumors recur upon drug discontinuation. Using *SKH1-Ptch1^{+/-}* as a model that closely mimics the spontaneous and accelerated growth pattern of BCCs in patients with BCNS, we show that AKT1, a serine/threonine

protein kinase, is intrinsically activated in keratinocytes derived from the skin of newborn *Ptch1^{+/-}* mice in the absence of carcinogenic stimuli. Introducing *Akt1* haplodeficiency in *Ptch1^{+/-}* mice (*Akt1^{+/-} Ptch1^{+/-}*) significantly abrogated BCC growth. Similarly, pharmacological inhibition of AKT with perifosine, an alkyl phospholipid AKT inhibitor, diminished the growth of spontaneous and UV-induced BCCs. Our data demonstrate an obligatory role for AKT1 in BCC growth, and targeting AKT may help reduce BCC tumor burden in BCNS patients. *Cancer Prev Res*; 9(10); 794–802. ©2016 AACR.

Introduction

Hedgehog (Hh) pathways are crucial for vertebrate fetal development (1). Of the three known mammalian Hh ligands, Sonic hedgehog (SHH) regulates segment polarity as well as a wide range of biological activities, from establishing left–right body symmetry and limb patterning to eye and central nervous system development. The other two Hh ligands, Desert hedgehog (DHH) and Indian hedgehog (IHH), are mainly involved in the development of male germ cells and cartilage, respectively (1). In the absence of SHH, *PTCH1*, a tumor suppressor and the Hh receptor, blocks Hh signaling by repressing a membrane-bound, G-protein–coupled, receptor-like protein known as smoothened (SMO) (1, 2). The binding of SHH to *PTCH1* relieves SMO repression, triggering a canonical Hh response whereby SMO moves to the primary cilium, and activates the GLI family of transcription factors (3). GLI forms a cytoplasmic complex with several accessory modulators, including the serine–threonine kinase Fused (FU), Suppressor of Fused (SUFU), and costal2 (COS2), a kinesin-related protein that binds the GLI-containing complex to microtubules. It is believed that SMO activity favors dissociation of these complexes and translocation of an active form of GLI from the cytoplasm to the nucleus, where it promotes the transcription

of Hh target genes, including *PTCH1*, *GLI*, *CCND1*, bone morphogenic proteins (BMPs), and a member of the TGF β superfamily (1, 2, 4).

The Hh pathway while highly active during human fetal development normally shuts down soon after birth. However, aberrant activation in adults drives the development and/or maintenance of numerous types of human malignancies, including cancers of the pancreas, prostate, and brain. It also regulates the proliferation of cancer stem cells (CSC), tumor progression, and metastases and may also hasten tumor relapse by augmenting multidrug resistance (MDR) pathways (2, 5). While most human tumors do not harbor somatic mutations in the Hh signaling pathway and demonstrate ligand-dependent Hh pathway activation (6), ligand-independent Hh pathway activation underlies the development of basal cell carcinoma (BCC), the most common type of human malignancy worldwide, due to loss-of-function mutations in *PTCH1*, gain-of-function mutations in *SMO*, as well as missense mutations in *GLI1* and *GLI3* (7). Germline mutations in *PTCH1* cause basal cell nevus syndrome (BCNS) or Gorlin syndrome (8). Individuals affected with BCNS typically develop large numbers of BCCs often beginning in early childhood and are at substantially increased risk for additional neoplasms such as medulloblastomas and rhabdomyosarcomas. The clinical utility of targeting aberrant Hh signaling in BCNS patients is illustrated by the results of our phase II clinical trial that showed that the orally administered SMO inhibitor vismodegib had remarkable efficacy in ablating BCCs (9). While these and other successes led to FDA approval of vismodegib for treating advanced/inoperable and metastatic BCCs, subsequent studies revealed significant tumor recurrence and acquired clinical resistance to vismodegib. This occurs during therapy primarily through secondary mutations in *SMO* (for example, D473G, D473Y, Q477E, and G497W) and, to a lesser extent, through concurrent copy number changes in *SUFU* and *GLI2*, thereby impairing drug binding and/or

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reactivating Hh pathways, confirming the importance of the Hh pathway in BCC growth (10–13). Several lines of evidence, however, demonstrate that additional pathways may synergistically contribute to BCC tumor growth. For example, there is a high risk of BCC in patients with cartilage-hair hypoplasia (CHH), an inherited disorder due to mutations in the *RMRP* (RNA component of mitochondrial RNA-processing endoribonuclease), and in patients with xeroderma pigmentosum (XP), an autosomal recessive disorder with defective nuclear excision DNA repair (14). Recently, a significant fraction of human BCCs was shown to carry mutations in various cancer-related genes that regulate a variety of cellular processes, including cell growth, differentiation, mitotic cycle, and oncogenic transformation (15). These data implicate involvement of multiple tumor driver pathways in BCC pathogenesis and suggest that identification of molecular targets distinct from SMO could help ameliorate the limitations of currently available SMO inhibitors and improve therapeutic index. One such molecular target that we have identified is AKT1, the serine/threonine protein kinase that regulates cell survival and is known to be dysregulated in numerous types of human cancer (16–20). Using SKH1-1-*Ptch1*^{+/−} mice that we recently described as a susceptible model for BCCs development, this study demonstrates that AKT1 is obligatory for BCC tumorigenesis, and that genetic and pharmacological inhibition of AKT prevents BCC growth.

Materials and Methods

Cells and reagents

Adult normal human epidermal keratinocytes (Lonza) and murine ASZ001 BCC cells (a gift from Dr. E. Epstein Jr.; ref. 21) were cultured in Medium 154CF, supplemented with Human Keratinocyte Growth Supplement (Thermo Fisher). ASZ001 cells display cellular morphology similar to that of human BCCs, express BCC markers, and are sensitive to SMO inhibition (22, 23). The authors did not authenticate the cell lines. pUSEamp-*myrAkt1* was obtained from Millipore. Perifosine was purchased from Selleckchem, edelfosine from Sigma-Aldrich, and MK-2206 from ChemieTek.

Western blotting, immunohistochemistry, and immunofluorescence

The experiments were performed as previously described (24–26). Antibodies against AKT1, AKT2, AKT3, p-AKT1 (S473), GLI1, and GLI2 were purchased from Cell Signaling Technology, SOX9 from Abcam, and β -actin from Sigma-Aldrich. Fluorescent images were acquired using the Zeiss LSM 5 Exciter confocal microscope with 40 \times oil immersion objective (Carl Zeiss) and analyzed using ImageJ software (NIH).

Proliferation assay

Proliferation was assessed using BrdUrd kit III (Roche Diagnostics Corp.), and Click-iT Plus EdU Alexa Fluor 488 Imaging Kit (Thermo Fisher) according to the manufacturers' instructions.

Colony formation assay

Primary human keratinocytes were transduced with the viral construct expressing *myrAkt1* (pUSEamp-*myrAkt1*) or pUSEamp vector (Millipore) at 10 μ g using lipofectamine 2000 (Invitrogen) and stable colonies were selected in the presence of G418 (Sigma-Aldrich). Virally transduced cells were then irradiated

with UV (30 mJ/cm²). Cells were then resuspended in top agar (0.7% agarose in keratinocyte media) and placed with the base layer prepared in 0.8% agar in keratinocyte media with supplements. After 3 weeks, cells were stained with Crystal Violet, and the number of colonies was counted. The experiment was repeated twice independently.

Generation of SKH1-*Ptch1*^{+/−} and SKH1-*Akt1*^{+/−} *Ptch1*^{+/−} mice and assessment of microscopic lesions

All animal experiments were performed in accordance with guidelines of our approved Columbia University Institutional Animal Care and Use Committee (IACUC) protocol. SKH1-*Ptch1*^{+/−} mice were generated by crossing B6-*Ptch1*^{+/−} (27) with SKH1 hairless mice (Charles River Laboratories). The resulting haired *Ptch1*^{+/−} F1 littermates were backcrossed for 10 generations to generate SKH1-*Ptch1*^{+/−} mice. SKH1-*Akt1*^{+/−} *Ptch1*^{+/−} mice were generated by crossing B6.129P2-Akt1tm1Mbb/J mice (Jackson Lab) with SKH1-*Ptch1*^{+/−} mice. The resulting haired *Akt1*^{+/−} *Ptch1*^{+/−} F1 littermates were backcrossed for 10 generations to generate SKH1-*Akt1*^{+/−} *Ptch1*^{+/−} mice. Tissue samples were prepared from H&E- and β -gal-stained full-thickness dorsal skin. For β -gal staining, glutaraldehyde and formalin-fixed tissues were treated with X-gal and iron buffer solution (Roche Diagnostics Corp.) for 48 hours and were processed using the manufacturer's protocol. Microscopic BCC-like lesions were defined as tumor islands composed of monomorphic basaloid cells with scant cytoplasm arranged as nests within the dermis. These lesions were counted as numbers per unit area and as total tumor area per square millimeter skin section as previously described (28). For each mouse, three skin strips (average 1.5 cm \times 0.1 cm) were analyzed.

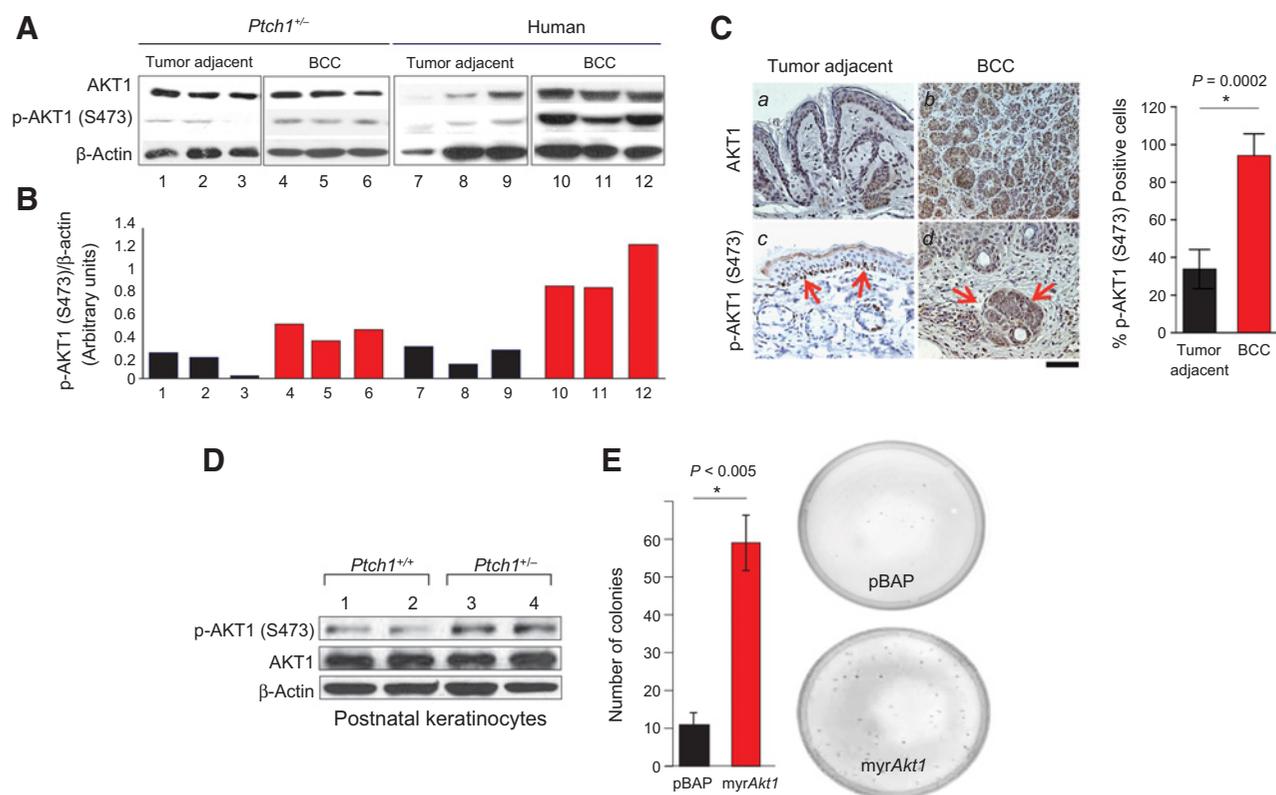
UV light source

A UV Irradiation Unit (Daavlin Co.) equipped with an electronic controller to regulate dosage was used. The UV source consisted of eight FS72T12-UV-HO lamps that emit UV (290–320 nm, 75%–80% of total energy) and UVA (320–380 nm, 20%–25% of total energy). A UVC sensor (Oriol's Goldilux UVC Probe) was routinely used during each exposure to confirm lack of UVC emission. The dose of UV was quantified with a UV Spectrum 305 Dosimeter obtained from the Daavlin Co. The radiation was further calibrated with an IL1700 Research Radiometer/Photometer from International Light Inc.

Tumor protocol in *Ptch1*^{+/−} and *Akt1*^{+/−} *Ptch1*^{+/−} mice

UV study: *Ptch1*^{+/−} and *Akt1*^{+/−} *Ptch1*^{+/−} and their wild-type *Ptch1*^{+/+} littermates were irradiated with UV (180 mJ/cm² twice per week for 30 weeks). **Itraconazole study:** *Ptch1*^{+/−} mice ($n = 20$) were UV-irradiated (180 mJ/cm² twice per week for 30 weeks) to induce BCCs. Irradiation was terminated. Mice were then treated with 2-hydroxypropyl- β -cyclodextrin ($n = 10$) or itraconazole ($n = 10$) (Sigma-Aldrich) (twice daily, 100 mg/kg, i.p.) for 24 days. The size of tumors was measured as previously described (28). For the perifosine study, control group received 0.9% NaCl/100 μ L (gavage) ($n = 8$), while the experimental group received perifosine (125 mg/kg body weight prepared in 0.9% NaCl, gavage, twice a week, $n = 8$) 30 minutes prior to each UV irradiation. All animals were irradiated with 180 mJ/cm² UV twice weekly for 32 weeks. Tumor number and tumor volume were recorded once a week and plotted in terms of weeks on test.

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**Figure 1.**

AKT1 is intrinsically activated in *Ptch1*^{+/-} keratinocytes and spontaneous BCCs. **A**, the levels of p-AKT1 (S473) in spontaneous murine BCCs developed in SKH1-*Ptch1*^{+/-} (*Ptch1*^{+/-}) mice and human BCCs. Western blotting assessed three representative BCCs from three different animals or three separate patients. β-Actin was used as an internal control. **B**, densitometric scanning of **A**. **C**, immunohistochemical assessment of AKT1 and p-AKT1 (S473) in tumor adjacent skin (a, c) and spontaneous BCCs (b, d) in *Ptch1*^{+/-} mice. Scale bar, 100 μm. Histogram represents percent p-AKT1 (S473)-positive cells in tumor-adjacent skins and BCCs in *Ptch1*^{+/-} mice. Three fields were counted. *, *P* < 0.05. **D**, p-AKT S473 levels are increased in primary keratinocytes isolated from postnatal day 2 *Ptch1*^{+/-} mice (lanes 3 and 4), but not in primary keratinocytes isolated from postnatal day 2 WT *Ptch1*^{+/+} mice (lanes 1 and 2). **E**, increased colony-forming capability of normal human keratinocytes expressing constitutively active AKT1 (pUSEamp-*myrAkt1*, *myrAkt1*). *, *P* < 0.005, compared with control. Normal primary human keratinocytes were transduced either with a viral construct expressing *myr-Akt1* (*myr-Akt1*) or vector (-, pBAP) only.

Statistical analyses

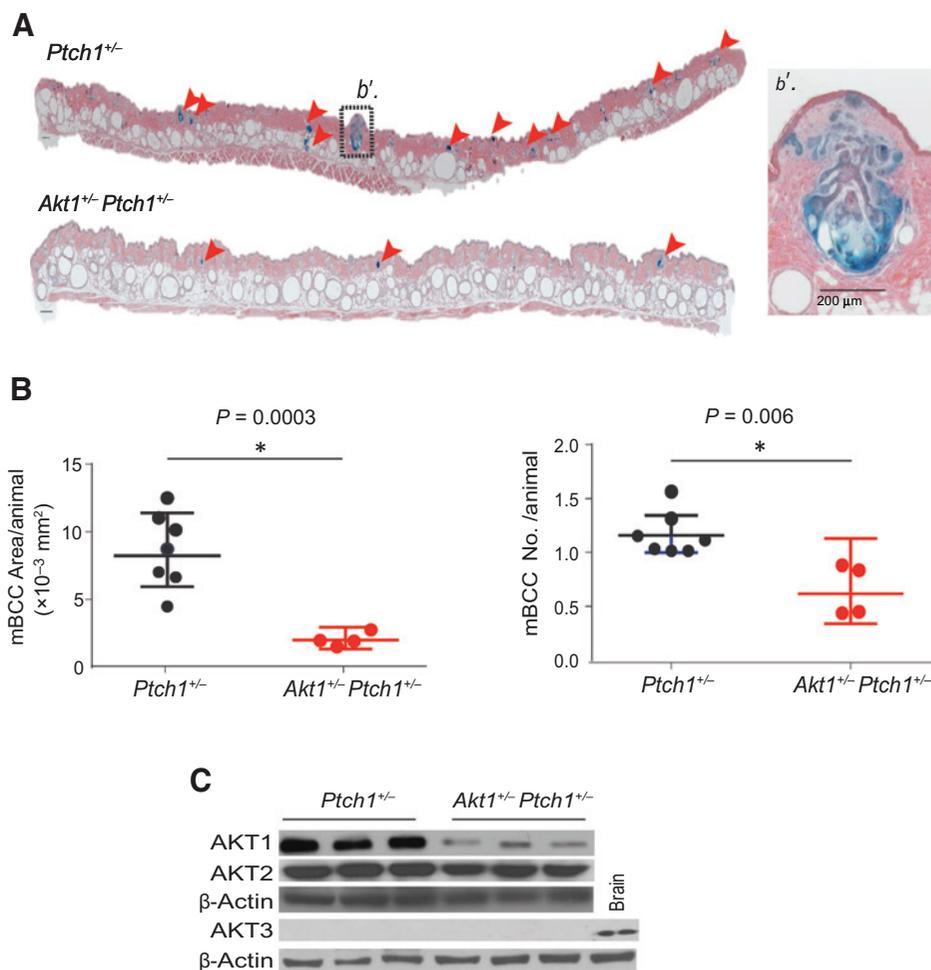
Statistical analyses were performed using the Student *t* test (two-tailed): *P* < 0.05 was considered statistically significant.

Results

AKT1 is intrinsically activated in *Ptch1*^{+/-} keratinocytes and spontaneous BCCs

We recently showed that the growth of BCC is greatly enhanced in hairless SKH1-*Ptch1*^{+/-} mice (26). Although the original introduction of *Ptch1*^{+/-} into the C57BL/6 strain (B6-*Ptch1*^{+/-}; ref. 27) rendered these animals susceptible to BCC following skin exposure to ionizing or UV radiation (29), these tumors grow quite slowly in these mice—perhaps due to their derivation from the C57BL/6 genetic background that is known to be inherently tumor resistant. In contrast, SKH1 hairless mice are highly susceptible to the growth of squamous cell carcinomas (SCC) following chronic UV exposure (25, 30). Chronic UV irradiation of SKH1-*Ptch1*^{+/+} wild-type (WT) mice results in the sequential growth of benign papillomas and SCCs in a pattern closely mimicking that of human SCCs, while these animals never

develop BCCs. By introducing *Ptch1* heterozygosity onto the tumor-susceptible SKH1 hairless background, we generated SKH1-*Ptch1*^{+/-} (hereafter referred to as *Ptch1*^{+/-}) mice. These animals exhibit the spontaneous growth of BCCs that characterizes patients with BCNS (31). AKT activation typically occurs in response to extracellular stimuli, and its phosphorylation at S473 is necessary for the full activation of the PI3K-AKT pathway (32). We detected S473 phosphorylation in BCCs that developed spontaneously in *Ptch1*^{+/-} mice and in human BCCs (Fig. 1A and B). AKT S473 phosphorylation was present in tumor cells but was undetectable in tumor stroma (Fig. 1C). Moreover, AKT S473 phosphorylation was detectable in primary keratinocytes isolated from newborn *Ptch1*^{+/-}, suggesting that AKT activation occurs in the absence of external stimuli in these cells (Fig. 1D). To confirm the role of AKT1 in cell survival and proliferation, we force activated AKT1 by transfecting myristoylated *Akt1* (*myr-Akt1*) into normal human primary keratinocytes. This resulted in enhanced colony formation (Fig. 1E). Together, these data suggest that the concurrent activation of both AKT and Hh pathways may play a crucial role in promoting BCC growth.

**Figure 2.**

Akt1 haploinsufficiency is sufficient to prevent the growth of spontaneous microscopic BCCs. **A**, representative H&E staining of skin sections from *Akt1*^{+/-} *Ptch1*^{+/-} mice and *Ptch1*^{+/-} littermates. Scale bar, 200 μ m.

B, assessment of size and number of spontaneous BCCs in *Akt1*^{+/-} *Ptch1*^{+/-} and *Ptch1*^{+/-}. Three skin strips (average 1.5 cm \times 0.1 cm) were analyzed for each mouse. Each dot represents data from one mouse. **C**, the levels of AKT isoforms in *Akt1*^{+/-} *Ptch1*^{+/-} mice and *Ptch1*^{+/-} littermates, assessed by Western blotting. 50 μ g protein per lane, actin as an internal control. Brain, brain extracts from *Akt1*^{+/-} *Ptch1*^{+/-}.

Genetic ablation of *Akt1* prevents the spontaneous growth of BCCs in *Akt1*^{+/-} *Ptch1*^{+/-} mice

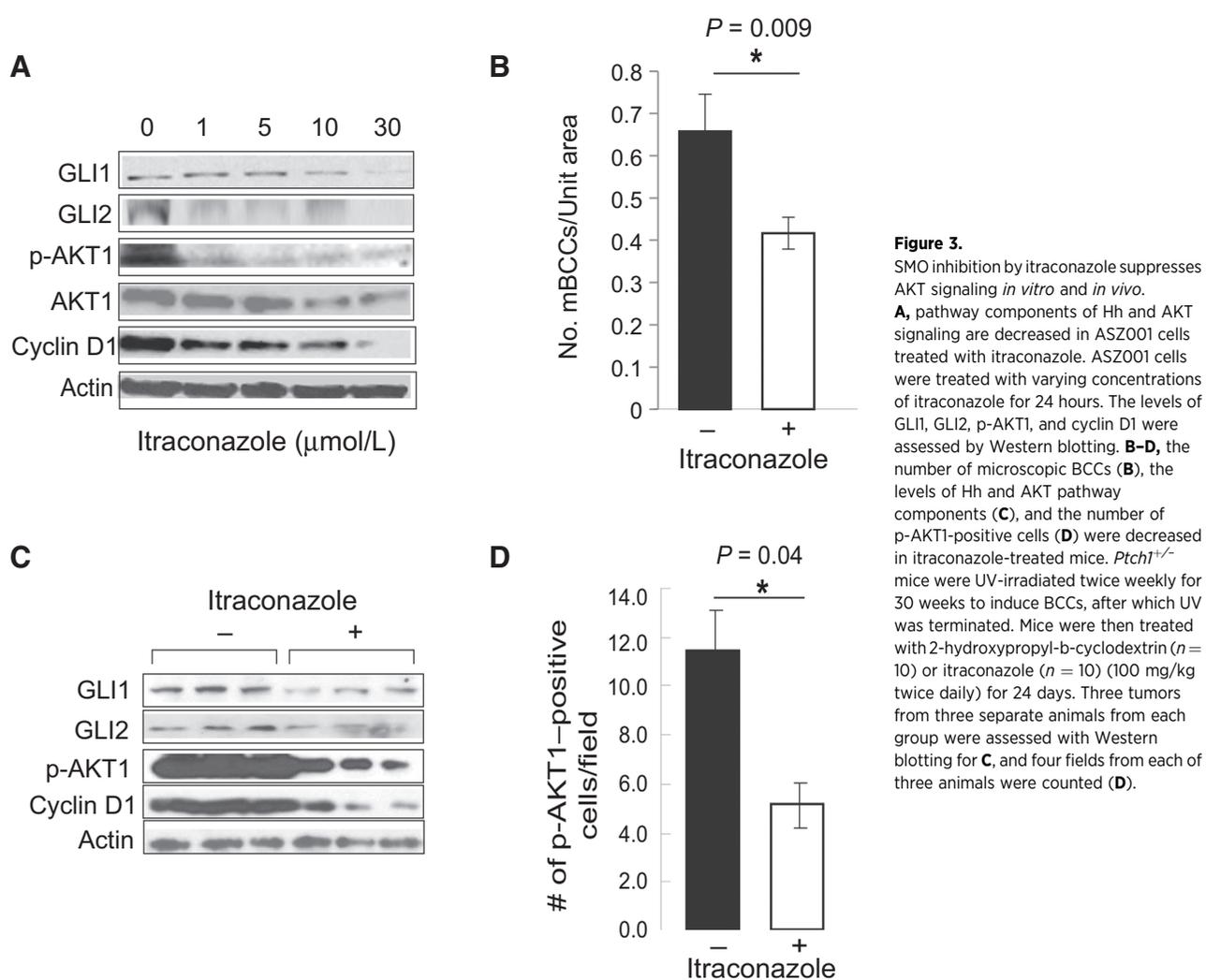
There are three known *Akt* isoforms, including *Akt1*, *Akt2*, and *Akt3*. *Akt1*-null mice manifest growth retardation (33), whereas *Akt2*-null mice display an insulin-resistant, diabetes-like syndrome (34), and *Akt3* null mice show impaired brain development (35). Because *Akt1* and *Akt2*, but not *Akt3*, are expressed in the skin, and because mouse embryo fibroblasts lacking *Akt1* (*Akt1*^{-/-} MEF) are sensitive to UV-induced apoptosis (33), we next assessed the effects of *Akt1* deletion in *Ptch1*^{+/-} mice. This study utilized *Akt1* haploinsufficient *Ptch1*^{+/-} mice, as *Akt1*^{-/-} littermates rarely survive. The few that do show severe growth retardation and high perinatal mortality (data not shown). Similar to BCNS patients where BCCs develop spontaneously even in sun-protected areas, spontaneous microscopic BCCs were detectable in our *Ptch1*^{+/-} mice starting as early as 8 weeks of age with 100% tumor incidence (ref. 26; Fig. 2). Figure 2A shows representative pictures of histological sections of skin from *Ptch1*^{+/-} and *Akt1*^{+/-} *Ptch1*^{+/-} mice at 12 months of age. *Ptch1*^{+/-} mice carry the insertion of a promoterless lacZ-neo fusion gene, thereby deleting a portion of exon 1 and all of exon 2 of *Ptch1*. Therefore, lacZ activity, mimicking an expression pattern of endogenous Hh target genes (e.g., *Ptch1*), serves as an accurate indicator of Hh pathway activation. In *Ptch1*^{+/-} mice, BCCs were detected by

β -gal staining (red arrowheads) with an inset showing a magnified view of an area with multiple BCCs (Fig. 2A, b'). *Akt1* haploinsufficiency resulted in substantial reductions in the tumor burden (in both size and number) of spontaneous BCCs in *Akt1*^{+/-} *Ptch1*^{+/-} mice as compared with their *Akt1* WT *Ptch1*^{+/-} littermates (Fig. 2B). Despite isoform-specific functions, it has been shown that *Akt1* knockdown can, in some instances, upregulate and activate AKT2, which in turn compensates for *Akt1* loss (36). Our analysis of *Akt1*^{+/-} *Ptch1*^{+/-} mice shows that AKT2 levels are no different than those in *Ptch1*^{+/-} mice (Fig. 2C), indicating non-overlapping roles for this AKT isoform and that AKT1 is crucial for the development of spontaneous BCCs in *Ptch1*^{+/-} mice.

SMO inhibition suppresses AKT1 signaling *in vitro* and *in vivo*

Itraconazole is an FDA-approved azole antifungal drug recently shown to be a potent, and specific inhibitor of Hh signaling (37). It is thought to reduce SMO translocation to the cilium (38). Treatment of murine ASZ001 cells (derived from BCCs induced in B6-*Ptch1*^{+/-}) with itraconazole (1–30 μ mol/L) dose dependently inhibited the growth of these cells (data not shown) and decreased the levels of the Hh components GLI1, GLI2, and cyclin D1 (Fig. 3A). Interestingly, itraconazole treatment decreased S473 phosphorylation (Fig. 3A). Cyclopamine, a natural compound known to block Hh signaling by binding to SMO, also suppressed

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**Figure 3.**SMO inhibition by itraconazole suppresses AKT signaling *in vitro* and *in vivo*.

A, pathway components of Hh and AKT signaling are decreased in ASZ001 cells treated with itraconazole. ASZ001 cells were treated with varying concentrations of itraconazole for 24 hours. The levels of GLI1, GLI2, p-AKT1, and cyclin D1 were assessed by Western blotting. **B–D**, the number of microscopic BCCs (**B**), the levels of Hh and AKT pathway components (**C**), and the number of p-AKT1-positive cells (**D**) were decreased in itraconazole-treated mice. *Ptch1*^{+/-} mice were UV-irradiated twice weekly for 30 weeks to induce BCCs, after which UV was terminated. Mice were then treated with 2-hydroxypropyl- β -cyclodextrin ($n = 10$) or itraconazole ($n = 10$) (100 mg/kg twice daily) for 24 days. Three tumors from three separate animals from each group were assessed with Western blotting for **C**, and four fields from each of three animals were counted (**D**).

AKT phosphorylation (data not shown). Oral administration of itraconazole (100 mg/kg twice daily for 24 days) suppressed the growth of existing UV-induced microscopic BCCs in *Ptch1*^{+/-} mice (Fig. 3B) and decreased Hh signaling, AKT phosphorylation, and the number of p-AKT-positive cells in tumors harvested from these animals (Fig. 3C and D). Taken together, these data indicate that AKT signaling is downstream of the Hh pathway.

Akt1 haploinsufficiency prevents UV-induced BCC growth in *Ptch1*^{+/-} mice

UV exposure is the major known risk factor for the induction of BCCs, both sporadically in the general population and in BCNS patients (39). Chronic UV irradiation of WT *Ptch1*^{+/+} mice induces the growth of SCCs (Fig. 4A, WT, red arrowheads; Fig. 4B), and these animals were largely resistant to the development of BCCs, as indicated by the absence of β -gal staining (Fig. 4A, WT). In *Ptch1*^{+/-} skin, chronic UV exposure resulted in extensive epidermal hyperplasia (*Ptch1*^{+/-} in Fig. 2A vs. Fig. 4A) and greatly enhanced the number and size of BCCs (Fig. 4A, *Ptch1*^{+/-}, black arrowheads). *Akt1* haploinsufficiency, however, substantially reduced the development of UV-induced BCCs in *Ptch1*^{+/-} mice (Fig. 4C). Extensive colocalization of p-AKT1 S473 with other

known BCC markers, GLI1 and SOX9, indicated their concurrent activation in *Ptch1*^{+/-} BCCs (Fig. 4D, *d* and *g*), compared with apparent reductions in AKT S473 phosphorylation and colocalization in *Akt1*^{+/-} *Ptch1*^{+/-} BCCs (Fig. 4D, *i*, *l*, and *o*). These data, together with substantial decreases in KI67, a cell proliferation marker, in *Akt1*^{+/-} *Ptch1*^{+/-} BCCs (Fig. 4D, *c* vs. *k*) further confirmed the importance of AKT1 in BCC growth.

Pharmacological inhibition of AKT inhibits UV-induced BCC growth

We next utilized a battery of AKT inhibitors that are currently in active clinical trials for various other human cancers to assess pharmacologic effects in ASZ001 murine BCC cells. These include the alkyl phospholipid perifosine and MK-2206, an allosteric inhibitor that binds to and inhibits AKT in a non-ATP competitive manner. Perifosine, in particular, has been shown to be relatively nontoxic and well tolerated in phase I/II clinical trials in patients with head and neck SCCs (40–42). Our results indicate that perifosine has a lower IC₅₀ for AKT (4 μ mol/L) in ASZ001 cells as compared with other inhibitors, which had IC₅₀ values ranging up to 50 μ mol/L (data not shown). Accordingly, we selected a concentration of 4 μ mol/L for further

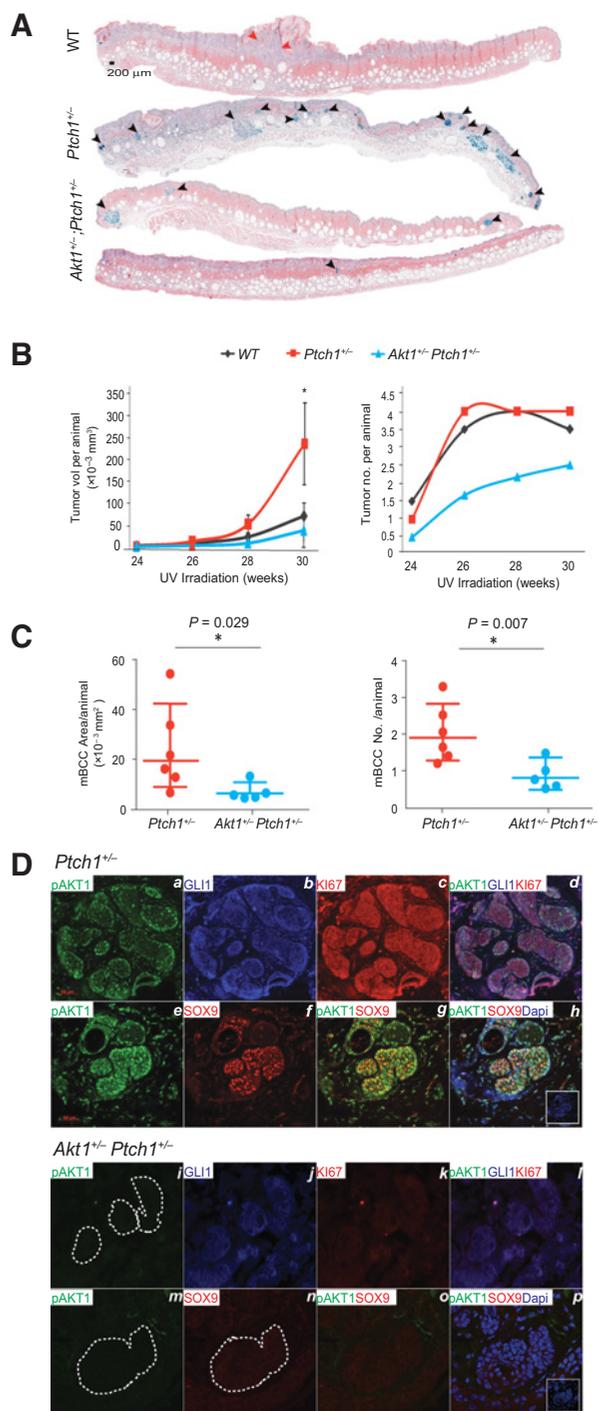


Figure 4. Haploinsufficient *Akt1* suppresses UV-induced BCC tumorigenesis in *Ptch1*^{+/-} mice. *Ptch1*^{+/-} ($n = 6$) and *Akt1*^{+/-};*Ptch1*^{+/-} ($n = 5$), and their WT littermates ($n = 6$) were irradiated with UV (180 mJ/cm², twice weekly) for 30 weeks. **A**, representative H&E staining of skin sections harvested at week 30. *Akt1* deficiency suppresses the growth of UV-induced skin tumors (**B**) and microscopic BCCs (**C**). Each dot in **C** represents data from one mouse. Three skin strips (avg. 1.5 cm \times 0.1 cm) were analyzed for each mouse. *, $P < 0.005$. **D**, colocalization of p-AKT with GLI1 and KI67, and with SOX9 in UV-induced BCCs in *Ptch1*^{+/-} (**a-h**) and *Akt1*^{+/-};*Ptch1*^{+/-} (**i-p**), as detected by immunofluorescence staining. Magnification, $\times 40$.

studies. Treatment of ASZ001 cells with perifosine, and its alkyl-lysophospholipid analogues, edelfosine, and MK-2206 inhibited AKT S473 and T308 phosphorylations (Fig. 5A). This was associated with substantially reduced proliferation of the ASZ001 cells as assessed by BrdUrd incorporation after 24 hours of treatment with 4 μ mol/L of each AKT inhibitor (Fig. 5B). More significantly, only perifosine and edelfosine induced apoptosis in ASZ001 cells, as shown by an increase in cleaved caspase-3 (Fig. 5A, left). Overexpression of myrAKT protected ASZ001 cells from perifosine-induced apoptosis, as shown by decreased levels of caspase-3 (Fig. 5A, right). While MK-2206 reduced p-AKT phosphorylation to levels comparable with that of perifosine or edelfosine (Fig. 5A), it did not affect the proliferative capacity of ASZ001 cells, at least up to a dose of 4 μ mol/L. Treatment with MK-2206 for 5 days still showed substantial amounts of EdU incorporation, indicative of cells in the S phase (Fig. 5C, g). In contrast, both perifosine and edelfosine completely abolished the proliferative capacity of these cells (Fig. 5C, e and f), and substantially decreased cell numbers (Fig. 5C, l and m). Perifosine appears to have superior efficacy in ASZ001 cells, as EdU-positive cells were barely detectable at day 2 (Fig. 5C, b). We, therefore, conducted a proof-of-concept preclinical investigation by administering perifosine orally (125 mg/kg, twice weekly) to chronically UV-irradiated *Ptch1*^{+/-} mice for 32 weeks. This regimen significantly reduced total tumor burden (Fig. 5D) as well as the size and number of microscopic BCCs (Fig. 5E). Our results demonstrate the potential utility of targeting AKT signaling to reduce tumor burden in human subjects who are at increased risk for BCC growth.

Discussion

In our previously published investigator-originated clinical trial, we showed that oral administration of the Hh pathway inhibitor vismodegib led to a dramatic reduction in the number of BCCs in BCNS patients, confirming the importance of Hh signaling in driving BCC pathogenesis. However, it soon became apparent in these patients that many of the BCCs that were no longer visible clinically and histologically rapidly recurred after drug discontinuation (9). These results while disappointing do raise intriguing questions regarding the source of the recurrent tumor cells and the mechanisms underlying their regrowth. Moreover, it is known that tumor resistance to vismodegib usually results from SMO mutations that block drug binding and/or reactivate Hh signaling (11, 12). These data collectively suggest that treatment with the currently available inhibitors of Hh signaling is unlikely to permanently eradicate BCCs, and that identification of additional tumor driver pathways distinct from the Hh pathway could lead to the development of innovative therapeutic approaches and could improve clinical outcomes.

Aberrant Akt activation is known to occur frequently in numerous types of human tumors, including those of the skin (16, 18, 19). BCNS patients develop a few to hundreds of BCCs, which in many cases grow aggressively, and may require multiple mutilating surgical procedures. In particular, spontaneous development of multiple BCCs in sun-protected skin is decidedly uncommon in the general human population, but is often a characteristic feature of BCNS patients. Except for the Hh pathway, the mechanisms underlying spontaneous tumor growth are largely unexplored, in

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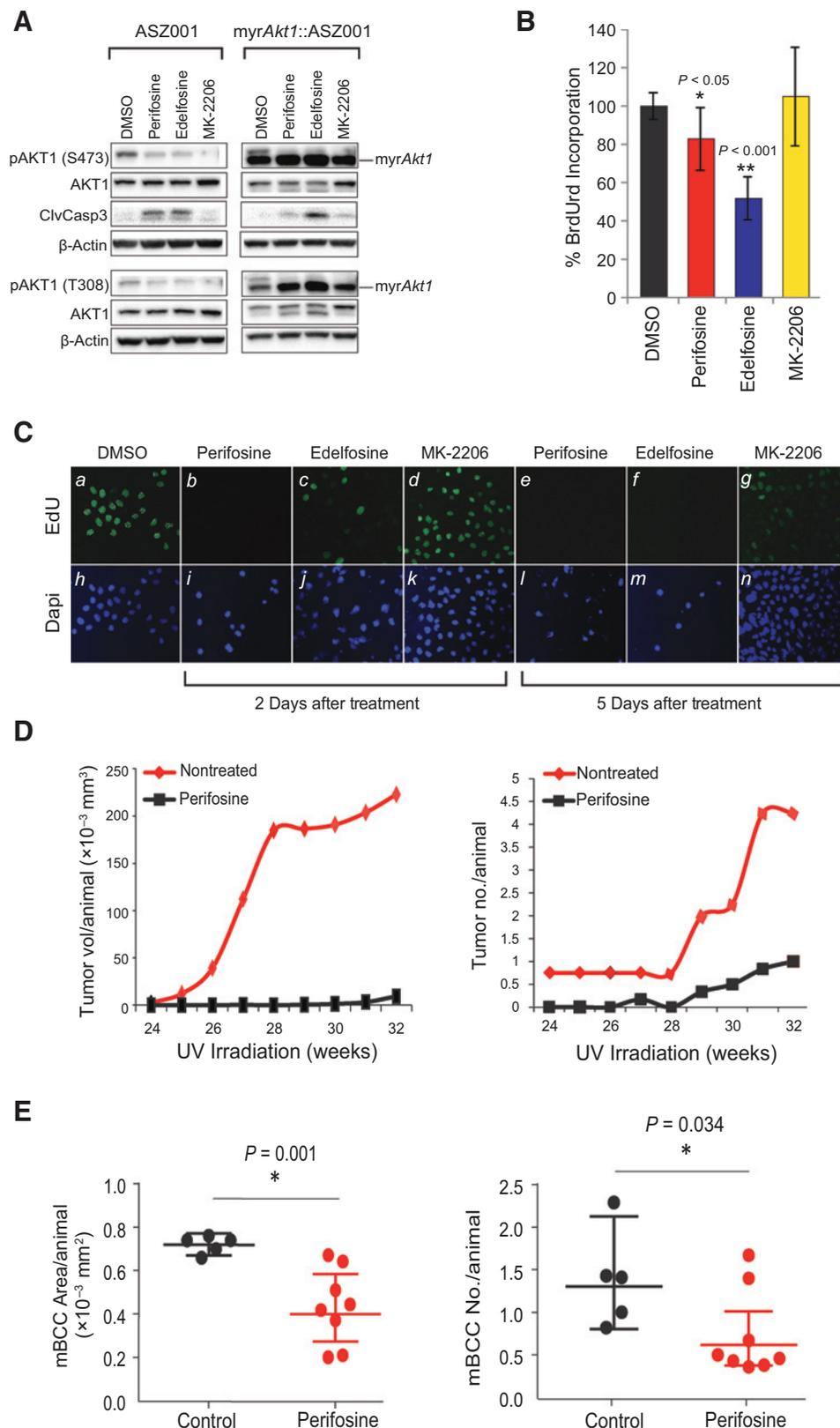


Figure 5.

Pharmacological inhibition of AKT signaling inhibits the proliferation of ASZ001 cells and suppresses UV-induced BCC growth. **A**, alkyl phospholipids reduce AKT phosphorylation and induce apoptosis in ASZ001 cells. ASZ001 cells were treated with 4 μmol/L of each AKT inhibitor, including alkyl phospholipids, and MK-2206, an allosteric AKT inhibitor, for 24 hours. The levels of AKT phosphorylation at S473 and T308 and cleaved caspase-3 were assessed by Western blotting. β-Actin serves as an internal control. **B** and **C**, AKT inhibition reduces proliferation of ASZ001 cells. Assessment of BrdUrd incorporation 24 hours (**B**) and EdU incorporation 2 or 5 days (**C**) after treatment with 4 μmol/L of each AKT inhibitor. Error bars, SD; *, $P < 0.05$; **, $P < 0.001$. Orally administered perifosine prevents the growth of UV-induced skin tumors (**D**) and microscopic BCCs (**E**) in *Ptch1*^{+/-} mice. Mice were irradiated with UV (180 mJ/cm², twice weekly) and received either 0.9% NaCl ($n = 8$) or perifosine (125 mg/kg of body weight, $n = 8$), twice weekly for 32 weeks. Each dot represents data from one mouse. Three skin strips (avg. 1.5 cm x 0.1 cm) were analyzed for each mouse.

part due to the lack of suitable animal models. SKH1-*Ptch1*^{+/-} mice are uniquely susceptible to the development of both spontaneous and UV-induced BCCs, thereby providing an animal model with features that closely resemble those of patients with BCNS. This study directly assessed the effects of *Akt1* loss by introducing *Akt1* haploinsufficiency in SKH1-*Ptch1*^{+/-} mice, and demonstrated that *Akt1* is essential for spontaneous BCC tumorigenesis. The absence of AKT S473 phosphorylation, indicative of the lack of its activity, was confirmed in *Akt1*^{+/-} *Ptch1*^{+/-} mice. The concomitant decreased levels of KI67, a marker of proliferation, and tumor burden in these mice further emphasize the importance of AKT1 function in cell survival and proliferation. UV and ionizing radiation are known to exacerbate BCC tumor burden in BCNS patients. The remarkable preclinical efficacy against UV-induced BCCs in *Ptch1*^{+/-} mice of the AKT inhibitor perifosine, selected over other inhibitors based on our *in vitro* studies, indicated that AKT could be an alternative viable target for reducing tumor burden in patients with BCNS.

Akt activation is posttranslational and is mediated through the upstream kinase cascade (e.g., phosphoinositide-dependent kinase 1 [PDK1]) in response to various growth factors and external stimuli. With respect to the Hh pathway, the activation of epidermal growth factor receptor (EGFR), platelet-derived growth factor receptor (PDGFR), and their downstream effectors (e.g., RAS/RAF/MEK/ERK and PI3K/AKT) has been shown to enhance GLI stability and transcriptional activity in various cancers (e.g., chronic lymphocytic leukemia, gastric cancer, and melanoma; refs. 43–47). AKT-mediated GLI activation has also been linked to drug resistance in breast cancer cells (48). While these data position AKT activation upstream of the Hh pathway, reciprocal regulation is also possible. Notably, GLI has been shown to transcriptionally regulate AKT1 expression in diffuse large B-cell lymphoma cells (49), and GLI-mediated AKT and c-MET phosphorylation was recently shown to promote migration and invasion of thyroid tumor cells (50). Our data, demonstrating *in vitro* and *in vivo* inhibition of AKT phosphorylation by the SMO inhibitor itraconazole, also imply that AKT acts downstream of Hh in BCC. In addition, despite the absence of external stimuli, AKT1 phosphorylation levels were significantly elevated in postnatal *Ptch1*^{+/-} keratinocytes compared with *Ptch1* WT keratinocytes, suggesting the existence of intrinsic mechanisms driving AKT phosphorylation. Elevated EGFR levels were previously reported in human BCCs, and forced expression of EGFR and GLI was shown to be critical for oncogenic transformation of spontaneously transformed, non-

tumorigenic HaCaT keratinocytes (51, 52). However, EGFR levels, as well as PDGFR, were barely detectable, and no apparent differences were observed in p-MEK1/2 levels, regardless of *Ptch1* status (data not shown). These results strongly suggest the lack of involvement of EGFR signaling in AKT phosphorylation in *Ptch1*^{+/-} keratinocytes; however, the exact mechanisms await further investigation. In addition, while GLI1 levels remained somewhat similar in UV-induced BCCs, irrespective of *Akt1* status, the apparent reduction in SOX9 levels in *Akt1*^{+/-} BCCs raises interesting questions with regard to the targets of AKT1 and its relevance to Hh signaling and BCC pathogenesis. Integration of the AKT and Hh pathways is undoubtedly under the influence of multiple regulatory pathways, which may account for the specificity and tumorigenic susceptibility of different target cells. Further investigation will likely reveal additional novel mechanism-driven targets for the chemoprevention and treatment of this most common type of human malignancy.

Disclosure of Potential Conflicts of Interest

D.R. Bickers is a consultant/advisory board member for SC Johnson Co. and has received an expert testimony from Goodwin Procter LLP. No potential conflicts of interest were disclosed by the other authors.

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Analysis and interpretation of data (e.g., statistical analysis, biostatistics, computational analysis): A.L. Kim, J.H. Back, Y. Zhu, N.P. Yardley, D.R. Bickers
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