

Spatially distinct roles of class Ia PI3K isoforms in the development and maintenance of PTEN hamartoma tumor syndrome

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PTEN hamartoma tumor syndrome (PHTS) comprises a collection of genetic disorders associated with germline mutations in the tumor suppressor gene *PTEN*. Therapeutic options and preventative measures for PHTS are limited. Using both genetically engineered mouse models and pharmacological PI3K isoform-selective inhibitors, we found that the roles of PI3K isoforms are spatially distinct in the skin: While p110 α is responsible for the sustained survival of suprabasal cells of the epidermis in the absence of PTEN, p110 β is important for the hyperproliferation of basal cells in PHTS. Furthermore, we identified a differential expression pattern of p110 α and p110 β in basal and suprabasal keratinocytes as well as differential PI3K regulation by upstream signals in the basal and suprabasal compartments of the epidermis, providing a potential molecular mechanism underlying the specific roles of PI3K isoforms in the epidermis. Finally, we demonstrate that combined inhibition of both PI3K isoforms prevents the development of PHTS and also reverses skin hamartomas that have reached advanced stages in mice. Together, these results not only advance our overall understanding of the diverse roles of PI3K isoforms, but also have the potential for meaningful translation via the clinical utilization of PI3K inhibitors for both prevention and therapy in PHTS patients.

[*Keywords:* genetic mouse models; PI3K inhibitors; PI3K isoforms; skin; PTEN hamartoma tumor syndrome]

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While somatic loss of *PTEN* tumor suppressor function is a frequent event in various sporadic human cancers (Sulis and Parsons 2003; Salmena et al. 2008), germline mutations in the *PTEN* gene cause several genetic or congenital disorders, including Cowden syndrome (CS), Bannayan-Ruvalcaba-Riley syndrome (BRRS), *Proteus* syndrome (PS) and *Proteus*-like syndrome (PLS), all characterized by multiple hamartomas occurring over various areas of the body. Collectively, these disorders are termed PTEN hamartoma tumor syndromes (PHTSs) (Blumenthal and Dennis 2008; Hobert and Eng 2009). CS, the prototypic syndrome, is characterized by a variety of skin lesions, which include papillomatous papules and acral and plantar keratosis, and is also associated with increased suscepti-

bility to breast, thyroid, and endometrial cancers (Liaw et al. 1997; Gustafson et al. 2007; Blumenthal and Dennis 2008; Hobert and Eng 2009). Notably, the occurrence of multiple skin hamartomas is the most common clinical manifestation of PHTS and occurs in almost all patients with CS (Hobert and Eng 2009), suggesting that PTEN/PI3K play important roles in the maintenance of skin epidermal homeostasis, as demonstrated by previous studies (Backman et al. 2004; Cully et al. 2006; Pankow et al. 2006).

PTEN is a lipid phosphatase that dephosphorylates the 3'-phosphoinositide products of the class Ia PI3Ks in epithelial cells by converting phosphatidylinositol (3,4,5) trisphosphate (PIP3) to phosphatidylinositol (4,5) bisphosphate (PIP2), thus counteracting class Ia PI3K activity. A reduction in 3'-phosphoinositides decreases the activity of kinases downstream from class Ia PI3Ks, such as Akt, and mTOR, and is responsible for much if not all of the

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tumor suppressor activity of PTEN (Sulis and Parsons 2003; Cully et al. 2006; Salmena et al. 2008). The class Ia PI3K family contains three highly homologous p110 catalytic isoforms (p110 α , p110 β , and p110 δ) encoded by three distinct genes: *Pik3ca*, *Pik3cb*, and *Pik3cd*, respectively. In mammals, the expression of p110 δ is largely restricted to the immune system, while p110 α and p110 β are expressed in all epithelial tissues and organs (Vivanco and Sawyers 2002; Liu et al. 2009). Recent studies have demonstrated that p110 α and p110 β play distinct roles in both signaling and cellular transformation (Jia et al. 2009; Vogt et al. 2009; Vanhaesebroeck et al. 2010). While p110 α is the major effector downstream from receptor tyrosine kinases (RTKs) (Zhao et al. 2006; Graupera et al. 2008; Utermark et al. 2012), p110 β is obligatory for signaling via G-protein-coupled receptors (GPCRs) and for RTK signaling in certain limited circumstances (Graupera et al. 2008; Guillermet-Guibert et al. 2008; Jia et al. 2008; Ciruolo et al. 2010). We previously reported that ablation of *p110 β* , but not *p110 α* , is sufficient to inhibit prostate intraepithelial neoplasia (PIN) induced by *Pten* loss in a mouse prostate tumor model (Jia et al. 2008). A recent study by Vanhaesebroeck's group (Berenjeno et al. 2012) using knock-in mouse models of kinase-dead alleles of *p110 α* or *p110 β* shows that, while, indeed, inactivation of *p110 β* but not *p110 α* blocks prostate tumor development, both *p110 α* and *p110 β* isoforms can modulate the impact of *Pten* loss in other tissues. Thus, identification of which p110 isoform is responsible for the transformation induced by *Pten* loss in any given tissue or tumor type remains intriguing as well as challenging.

Results

Genetic ablation of both p110 α and p110 β isoforms of PI3K completely blocks the development of multiple skin hamartomas induced by Pten loss

We set out to identify the specific roles of the *p110 α* and *p110 β* isoforms of class Ia PI3K in a murine model of CS. We first crossed mice carrying a floxed *Pten* allele with mice carrying a *K14Cre* transgene, as described in a previously reported mouse model of CS (Squarize et al. 2008). Over time, these mice developed multiple dermal lesions closely resembling skin PHTS. We then generated cohorts of compound mice with *K14Cre*, *K14Cre;Pten^{L/L}*, *K14Cre;Pten^{L/L};p110 α ^{L/L}*, *K14Cre;Pten^{L/L};p110 β ^{L/L}*, and *K14Cre;Pten^{L/L};p110 α ^{L/L};p110 β ^{L/L}* genotypes (hereafter termed wild-type, *Pten^Δ*, *Pten^Δ;p110 α ^Δ*, *Pten^Δ;p110 β ^Δ*, and *Pten^Δ;p110 α ^Δ;p110 β ^Δ* mice) to facilitate concurrent ablation of *Pten* with either *p110 α* or *p110 β* in epidermal keratinocytes. These mice were viable and appeared normal from birth up to 3 wk of age. The *Pten^Δ* mice developed skin lesions progressively after weaning and, within 12 wk, presented with multiple cutaneous hamartomas found all over the body. The most visible symptoms were papillomatous lesions around the facial orifices, ears, and paws, and most of these were associated with hyperkeratosis (Fig. 1A; Supplemental Fig. 1A). The progression of skin hamartoma in *Pten^Δ* mice was eval-

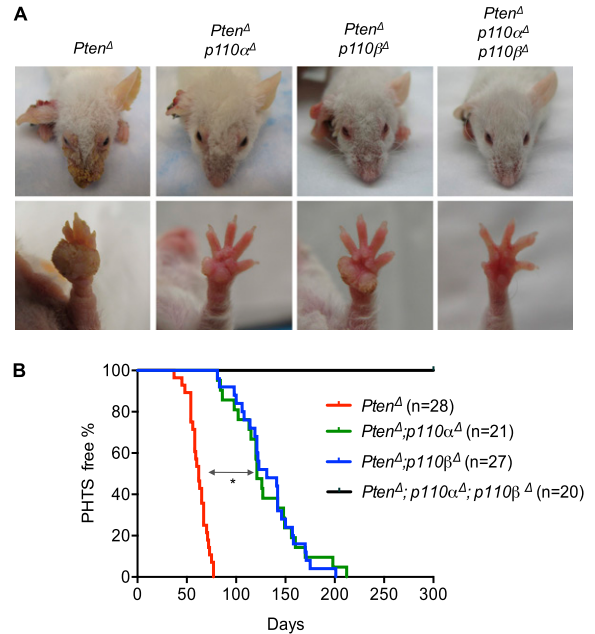


Figure 1. Effects of genetic ablation of *p110 α* and/or *p110 β* on the development of PHTS in mice. (A) Representative photos of the heads and left front paws of mice of the indicated genotypes at 12 wk of age. (Δ) Deletion status of the floxed alleles. (B) Kaplan-Meier (KM) curves showing the onset of PHTS in mice of the indicated genotypes. The median skin hamartoma onset for *Pten^Δ* mice (red line) is 62 d. Ablation of either *p110 α* (green line) or *p110 β* (blue line) delayed symptom onset to 121 and 131 d, respectively. (*) $P < 0.001$ (log-rank test). All *Pten^Δ;p110 α ^Δ;p110 β ^Δ* mice remained free of PHTS symptoms for at least 300 d (black line).

uated according to the pathological criteria of PHTS as described in previous studies (Supplemental Fig. 1B; Nelen et al. 1996; Eng 2000; Laury et al. 2011). The median onset of disease in *Pten^Δ* mice was 62 d (Fig. 1B, red line). There were no significant differences in the severity and time to onset of these skin lesions comparing male and female *Pten^Δ* mice (Supplemental Fig. 2). Although ablation of either the *p110 α* or *p110 β* gene significantly delayed the development and severity of the skin lesions induced by *Pten* loss in keratinocytes (Fig. 1; Supplemental Fig. 1B), all *Pten^Δ;p110 α ^Δ* and *Pten^Δ;p110 β ^Δ* mice developed skin hamartomas with a median latency of 121 and 131 d, respectively (Fig. 1B). Remarkably, however, *Pten^Δ;p110 α ^Δ;p110 β ^Δ* mice, in which both *p110 α* and *p110 β* isoforms were ablated, did not develop hamartomas over an observation period of 300 d (Fig. 1). This finding suggests that both p110 α and p110 β are essential to the development of skin lesions in the absence of *Pten*.

Loss of both p110 α and p110 β isoforms of PI3K is required to counteract Pten loss-induced hyperphosphorylation of Akt and restore the normal skin thickness and architecture in the absence of Pten

To examine the effect of the PI3K/*Pten* pathway activation on the epidermis of adult mice, we carried out

histological analyses on skin biopsies from adult mice with various PI3K/*Pten* genotypes at 8 wk of age, prior to the formation of hamartomas. *Pten*^Δ mice showed significantly thickened skin with a marked increase in the number of epidermal cell layers (Fig. 2A). Keratinocyte-specific *Pten*-null mice with concomitant ablation of either PI3K isoform (*Pten*^Δ;*p110α*^Δ or *Pten*^Δ;*p110β*^Δ mice) did not display increased skin thickness to the same degree as the *Pten*^Δ mice, but their epidermal layers were still significantly thicker than those of wild-type control

mice (Fig. 2A). Notably, in *Pten*-deficient epidermis, concurrent ablation of both PI3K isoforms (*Pten*^Δ;*p110α*^Δ;*p110β*^Δ) restored epidermal thickness and skin architecture to a normal state (Fig. 2A). Consistent with this, Western blot analyses of epidermal samples isolated from mice with these genotypes demonstrated that the increased levels of Akt phosphorylation observed in the absence of *Pten* were partially reduced by ablation of either *p110α* or *p110β* (Fig. 2B). Only concurrent deletion of both *p110α* and *p110β* reduced the elevated Akt phosphorylation

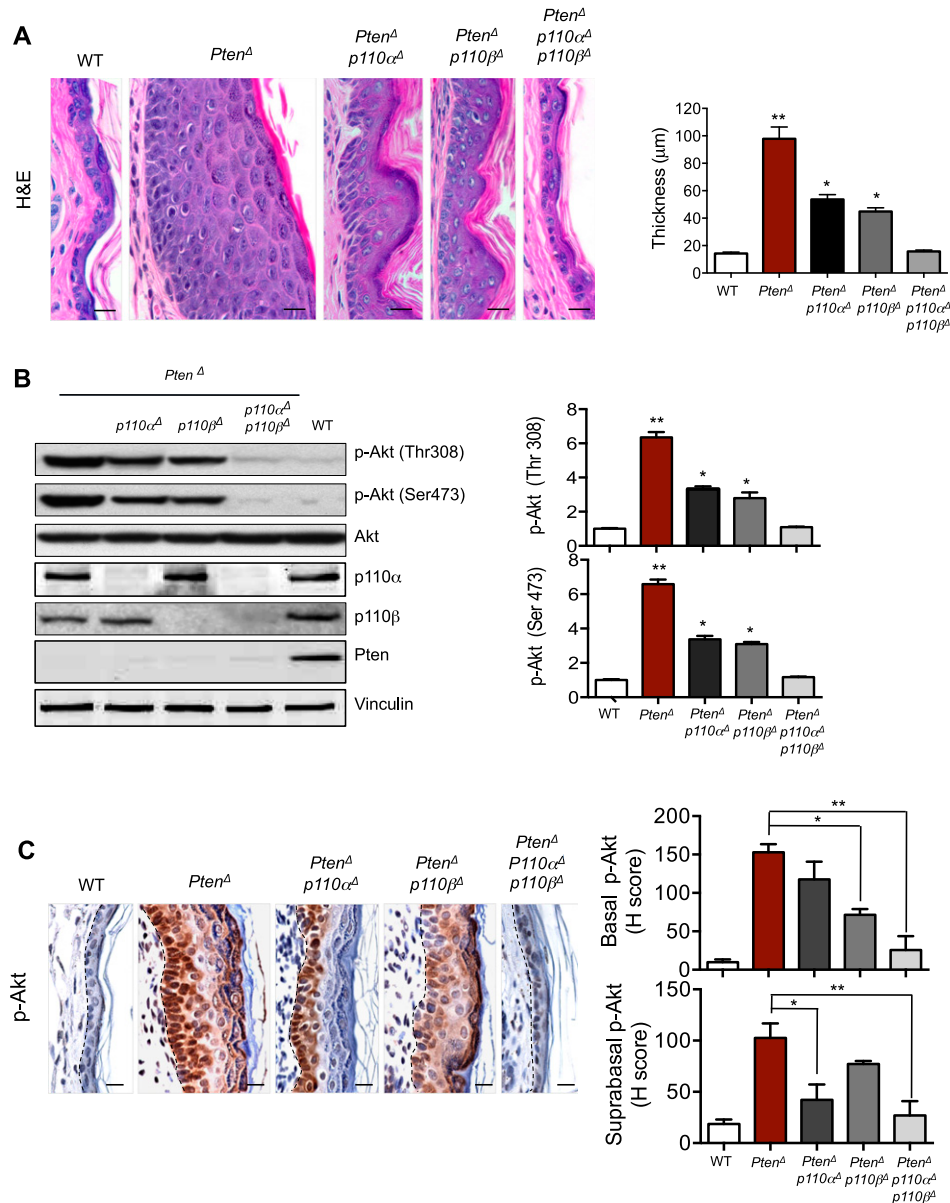


Figure 2. Effects of *p110α* and/or *p110β* ablation on skin epidermis in the absence of *Pten*. (A) hematoxylin and eosin (H&E) staining of skin sections from ears of 8-wk-old mice of the indicated genotypes. Bar, 20 μm. The bar graph represents the thickness of the epidermis from mouse ears of the indicated genotypes. Data are shown as mean ± SEM ($n = 6$ for each genotype). (*) $P < 0.01$; (**) $P < 0.001$ (Student's *t*-test). (B) Western blot analyses of p-Akt in primary keratinocytes isolated from mice of the indicated genotypes. Data are shown as mean ± SEM ($n = 6$). (*) $P < 0.01$; (**) $P < 0.001$ (Student's *t*-test). (C) IHC analyses of p-Akt in epidermal sections of ears from 8-wk-old mice of the indicated genotypes. Dashed lines mark the epidermal–dermal junction. Bar, 20 μm. Histogram scores (H-scores) of p-Akt in basal and suprabasal epidermis are shown as mean ± SEM ($n = 6$). (*) $P < 0.01$; (**) $P < 0.001$ (Student's *t*-test).

lation levels induced by *Pten* loss to normal levels (Fig. 2B).

Immunohistochemistry (IHC) analysis of phospho-Akt (p-Akt) revealed compartmentalization differences not detectable in Western blotting: While cytoplasmic p-Akt levels were increased in cells in all layers of the thickened *Pten*-null skin, surprisingly, we found nuclear p-Akt staining in cells in the inner layers of the epidermis, forming two zones of p-Akt staining (Fig. 2C; Supplemental Fig. 3). More interestingly, while ablation of *p110 α* markedly reduced the cytoplasmic p-Akt signal in the outer layer cells of the *Pten*-null skin, ablation of *p110 β* greatly diminished nuclear p-Akt in the inner layer cells of *Pten*-null skin (Fig. 2C; Supplemental Fig. 3). The finding of nuclear p-Akt in basal cells of *Pten*-null epidermis is reminiscent of nuclear p-Akt found in neoplastic colon and prostate tissues in *Pten*^{+/-} *Pml*^{-/-} mice (Trotman et al. 2006). While the mechanism underlying this observation remains to be determined, our finding delineates additional isoform-specific roles and, further, suggests that *p110 α* and *p110 β* isoforms play differential roles in spatially defined portions of the adult skin epidermis.

The p110 α isoform is responsible for the sustained survival of suprabasal cells of epidermis in the absence of Pten, while p110 β is important for the hyperproliferation of basal cells in epidermal PHTS

Mammalian skin epidermis is composed of stratified epithelium with an underlying basement membrane, a single layer of proliferating basal cells, and layers of suprabasal cells that undergo differentiation in step-wise stages progressing from the spinous layer, granular layer, and outermost stratum corneum (Fuchs and Nowak 2008). Normal adult skin maintains epidermal homeostasis by balancing the proliferation, differentiation, and apoptosis of epidermal cells in these layers (Blanpain and Fuchs 2009). To examine the divergent effects of PI3K isoforms on *Pten*-null skin, we further analyzed skin biopsies from mice with various *p110/Pten* genotypes at 8 wk of age, as described above. Keratins 5/14 (K5/K14) and K1/K10 are protein markers commonly used to indicate basal and suprabasal cells, respectively. However, in *Pten*-null epidermis, K5 is expressed in almost the entire thickened epidermis and overlaps with K10 in the suprabasal layers, indicating a defective transition in the expression of these structural proteins in the basal-to-suprabasal shift, when *Pten* is lost in the epidermis (Fig. 3A). Δ Np63, the predominant p63 isoform expressed in mature epidermis, is preferentially expressed in basal epidermis (Yang et al. 1998; Liefer et al. 2000). Immunostaining analysis revealed that Δ Np63 remains as the predominant p63 isoform in *Pten*-null epidermis (Supplemental Fig. 4A). Notably, in the *Pten*-null epidermis, unlike the massive expression pattern of K5, Δ Np63-positive cells display relatively modest expansion into the suprabasal space (Supplemental Fig. 4A,B). Moreover, Δ Np63 and K10 expressions show a discrete separation in the epidermis (Supplemental Fig. 5). Many cells in these Δ Np63-positive layers are also Ki67- and BrdU-positive

(Fig. 3B,C; Supplemental Fig. 6), indicating that, in the absence of *Pten*, Δ Np63 remains as a reliable marker for basal cells with increased proliferation capacity. Consistent with the expanded suprabasal layers of *Pten*-null skin as observed through K10 staining, a marked accumulation of loricrin-positive (a marker for terminally differentiated epidermal cells) cells was also found in the granular layer of the *Pten*-deficient skin (Fig. 3D).

Interestingly, ablation of *p110 α* in *Pten*-null skin largely prevented the accumulation of K10- and loricrin-positive cells but did not significantly affect the distribution of Ki67- and Δ Np63-positive cells (Fig. 3A–D). Conversely, deletion of *p110 β* in *Pten*-null epidermis greatly reduced K5-positive cells and confined both Ki67- and p63-positive cells to the basal layer but still permitted abnormal accumulation of the terminally differentiated granular cells (Fig. 3A–D). Concurrent ablation of both *p110 α* and *p110 β* isoforms restored epidermal thickness, proliferation indexes, and skin architecture in *Pten*-deficient epidermis to a normal state (Fig. 3A–D). Together, these findings suggest that *p110 α* is primarily responsible for the survival of differentiated suprabasal cells, while *p110 β* is important for the proliferation of basal cells.

To further understand the unique, spatially distinct roles of the PI3K isoforms in the epidermis, we determined the expression pattern of these two isoforms in basal and suprabasal cells. The expression of α 6-integrin is restricted to basal cells and has been used to sort basal and suprabasal keratinocytes (Yi et al. 2008). We also confirmed that α 6-integrin expression is largely limited in Δ Np63-positive basal layers in *Pten* ^{Δ} epidermis. We therefore isolated skin keratinocytes and collected basal cells (α 6-integrin high) and suprabasal cells (α 6-integrin low) through fluorescence-activated cell sorting (FACS) (Supplemental Fig. 7). Real-time quantitative PCR of these cells isolated from wild-type skin revealed that the expression level of *p110 α* in suprabasal cells is approximately fivefold higher than that of basal cells. Conversely, the expression level of *p110 β* in basal cells is approximately fivefold higher than in suprabasal cells (Fig. 3E). The cells isolated from the *Pten*-null skin show the same differential expression pattern of *p110 α* and *p110 β* in basal and suprabasal cells of the epidermis (Fig. 3F). The differences were smaller than those observed in wild-type skin, perhaps due to the perturbed basal-to-suprabasal transition in the absence of *Pten*, leading to a possible contamination of basal cells with suprabasal cells and vice versa during cell sorting. These data suggest that the differential expression of *p110 α* and *p110 β* in the basal and suprabasal compartments of epidermis is likely one of the mechanisms responsible for the spatially differential roles of these two *p110* isoforms in skin epidermis.

A p110 α -selective inhibitor (A66) induces cell death in the surface layers of epidermis, while a p110 β -selective inhibitor (KIN-193) reduces the proliferation of basal cells in skin PHTS

To further define the distinct roles of PI3K isoforms in established skin hamartomas, we extended our study to

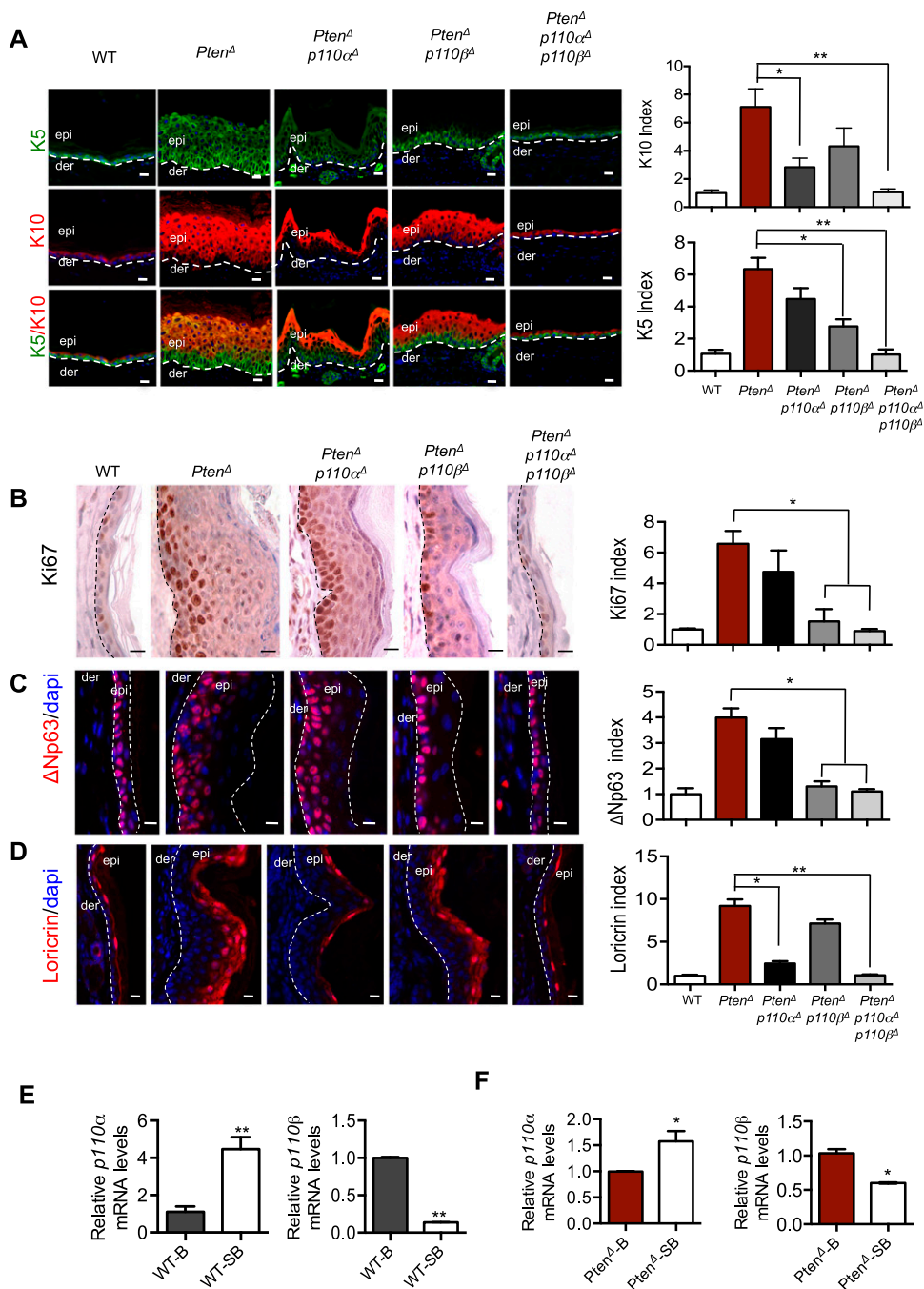


Figure 3. Divergent roles of PI3K isoforms on the skin of *Pten*^Δ mice. (A) Immunofluorescence (IF) staining analyses of K5 and K10 in epidermal sections from the ears of 8-wk-old mice of the indicated genotypes. Dashed lines denote the dermo/epidermal border. Bar, 20 μ m. The number of K5- and K10-positive cells in a 100- μ m length of epidermis of wild-type mice was used as a control (defined as 1). The bar graph represents the K5 or K10 index of each genotype relative to the control (\pm SEM; $n = 6$ for each genotype). (*) $P < 0.01$; (**) $P < 0.001$ (Student's *t*-test). (B) IHC analyses of Ki67 in epidermal sections of ears from 8-wk-old mice of the indicated genotypes. Dashed lines denote the dermo/epidermal border. Bar, 20 μ m. The number of Ki67-positive cells in a 100- μ m length of epidermis of wild-type mice was used as a control (defined as 1). The bar graph represents the Ki67 index of each genotype relative to the control (\pm SEM; $n = 6$ for each genotype). (*) $P < 0.01$ (Student's *t*-test). (C) IF staining analyses of the basal marker Δ Np63 in epidermal sections of ears from 8-wk-old mice of the indicated genotypes. (der) Dermis; (epi) epidermis. The area between the two dashed lines denotes the epidermal layer. Bar, 20 μ m. The number of Δ Np63-positive cells in a 100- μ m length of epidermis of wild-type mice was used as a control (defined as 1). The bar graph represents the Δ Np63 index of each genotype relative to the control (\pm SEM; $n = 6$ for each genotype). (*) $P < 0.01$ (Student's *t*-test). (D) IF staining analyses of the terminal differentiation marker loricrin in epidermal sections of ears from 8-wk-old mice of the indicated genotypes. Dashed lines denote the dermo/epidermal border. Bar, 20 μ m. The number of loricrin-positive cells in a 100- μ m length of epidermis of wild-type mice was used as a control (defined as 1). The bar graph represents the loricrin index of each genotype relative to the control (\pm SEM; $n = 6$ for each genotype). (*) $P < 0.01$; (**) $P < 0.001$ (Student's *t*-test). (E) Quantitative RT-PCR of *p110*^α and *p110*^β in FACS-purified cells from ear epidermis of 8-wk wild-type mice. (B) Basal layer; (SB) suprabasal. The relative expression level of each gene is normalized against its basal cell expression, defined as 1. (*) $P < 0.01$; (**) $P < 0.001$ (Student's *t*-test). (F) Quantitative RT-PCR of *p110*^α and *p110*^β in FACS-purified cells from ear epidermis of 8-wk *Pten*^Δ mice. (B) Basal layer; (SB) suprabasal. The relative expression level of each gene is normalized against its basal cell expression, defined as 1. (*) $P < 0.01$; (**) $P < 0.001$ (Student's *t*-test).

pharmacological approaches, isolating skin biopsies from hamartoma-bearing *Pten*^A mice treated for 3 d with vehicle, BKM120 (a pan PI3K inhibitor) (Maira et al. 2012), A66 (a p110 α -selective inhibitor) (Jamieson et al. 2011), and KIN-193 (a p110 β -selective inhibitor) (Ni et al. 2012). Analyses of skin sections demonstrated that, while A66 significantly reduced the number of K10-positive supra-basal cells, KIN-193 greatly reduced the number of Δ Np63-positive basal cells (Fig. 4A). Notably, BKM120 treatment reduced both K10- and Δ Np63-positive cells in the thickened *Pten*-null epidermis (Fig. 4A). Further analyses of these skin sections showed that, while A66 eliminated the majority of p-Akt in the outer layer cells, KIN-193 reduced p-Akt—in particular, nuclear p-Akt—in basal layer cells, and BKM120 almost completely abolished p-Akt in *Pten*-null skin (Fig. 4B; Supplemental Fig. 8), mirroring the genetic findings described above. Moreover, we found that A66 selectively induced cell death of loricrin-positive granular cells but had little effect on the proliferative capacity of the basal layer cells (Fig. 4C,D; Supplemental Fig. 9). In contrast, KIN-193 significantly reduced the number of Ki67-positive cells in basal layers but failed to induce cell death of granular cells (Fig. 4C,D; Supplemental Fig. 9). Consistently, BKM120 treatment markedly reduced Ki67-positive cells in the basal layer and greatly increased numbers of TUNEL-positive cells in the granular layer of epidermis as compared with those from vehicle controls (Fig. 4C,4D; Supplemental Fig. 9). Together, these pharmacological results are consistent with our genetic data and further support the spatially differential roles of p110 α and p110 β in skin epidermis.

Previous studies have demonstrated that the p110 α isoform mediates RTK signaling, and the p110 β isoform engages GPCR signaling (Vanhaesebroeck et al. 1997; Zhao et al. 2005; Jia et al. 2008; Kroeze et al. 2012; Utermark et al. 2012). We therefore sought to determine whether the specific roles of p110 isoforms in epidermal-lacing *Pten* are differentially regulated by upstream signals that mediate the activation of PI3K, e.g., growth factors and chemokines. We treated hamartoma-bearing mice with RTK inhibitors (a combination of lapatinib and sunitinib to achieve inhibition of a broad spectrum of RTKs) or a general GPCR signaling inhibitor (pertussis toxin) and harvested the skin specimens for analysis 3 d after treatment. IHC staining for p-Akt revealed that the treatment with RTK inhibitors reduced p-Akt in supra-basal cells, while the treatment with the GPCR signaling inhibitor reduced p-Akt in basal layer cells (Fig. 4E). Moreover, we showed that RTK inhibitors induced apoptosis of granular cells, whereas GPCR inhibitor suppressed proliferation of basal cells (Fig. 4F,G). In addition, we found that cells expressing CXCR3 (a member of the GPCR family) are tightly clustered with basal cells that express high α 6-integrin in both wild-type and *Pten*-null skin (Supplemental Fig. 10), consistent with a previous finding that CXCR3 is highly enriched in basal cells (Kroeze et al. 2012). Together, the differences in the upstream regulation of the PI3K signaling in different regions of the epidermis may act in concert with the differential expressions of the two isoforms in the skin

layers, providing a potential molecular mechanism underlying the spatially distinct roles of p110 α and p110 β isoforms in the epidermis.

A pan-PI3K inhibitor, BKM120, is able to prevent the development of skin PHTS

Since our genetic studies suggest that uncontrolled PI3K activity of both isoforms is responsible for the development of skin hamartomas in the absence of *Pten*, we hypothesized that pharmacological inhibition of class Ia PI3K activity might interfere with the initiation of PHTS in mice. We therefore tested the effects of BKM120, a pan-class I PI3K inhibitor currently in clinical trials for cancer therapy (Maira et al. 2012). We treated *Pten*^A mice at 3 wk of age with various doses of BKM120 and isolated epidermal tissues for Western blot analysis of p-Akt to assess the inhibitory effects of this drug. A relatively low dose of BKM120 (25 mg/kg) reduced p-Akt levels in keratinocytes of *Pten*^A mice to baseline levels (Supplemental Fig. 11). We then subjected cohorts of *Pten*^A mice to daily treatments with 25 mg/kg BKM120 ($n = 14$) or vehicle ($n = 12$) beginning at 3 wk of age. As expected, all mice in the vehicle control group progressively developed various skin lesions within 60 d (Fig. 5A; Supplemental Fig. 12). Remarkably, all mice in the treatment group were free of skin lesions during the entire 11-wk course of treatment (Fig. 5; Supplemental Fig. 12). However, when BKM120 was withdrawn from six mice in the BKM120 treatment group at the end of the 11th week, all six developed skin lesions within a week (Fig. 5; Supplemental Fig. 12). In contrast, the remaining eight mice in the cohort, which were maintained on BKM120 treatment, remained symptom-free (Fig. 5A; Supplemental Fig. 12), suggesting that continuous drug treatment is required to prevent the development of PHTS. Accordingly, histological analyses of skin samples from the faces and paws of mice at the end of the treatment course demonstrated that normal skin thickness and architecture were maintained in mice treated with BKM120 (Fig. 5C). Our data indicate that a submaximal and well-tolerated dose of a pan-class Ia PI3K inhibitor can offset the excessive proliferation of epidermal cells and prevent the survival/accumulation of terminally differentiated epidermal cells in the absence of *Pten*, thereby maintaining normal skin morphology and minimizing the phenotypic consequences of *Pten* deficiency.

BKM120 is able to reverse skin PHTS that has reached advanced stages in mice

Since inhibition of p110 α promotes cell death of accumulated outer layer epidermal cells and inhibition of p110 β suppresses hyperproliferation of inner layer basal cells on PHTS skin, we hypothesized that combined inactivation of both p110 α and p110 β using a pan-PI3K inhibitor might induce regression of hamartomas in *Pten*-deficient epidermis and alleviate disease symptoms in adult mice with established PHTS. Using *Pten*^A mice that had developed advanced multiple skin hamartomas, a clinically relevant dose of BKM120 (45 mg/kg) was

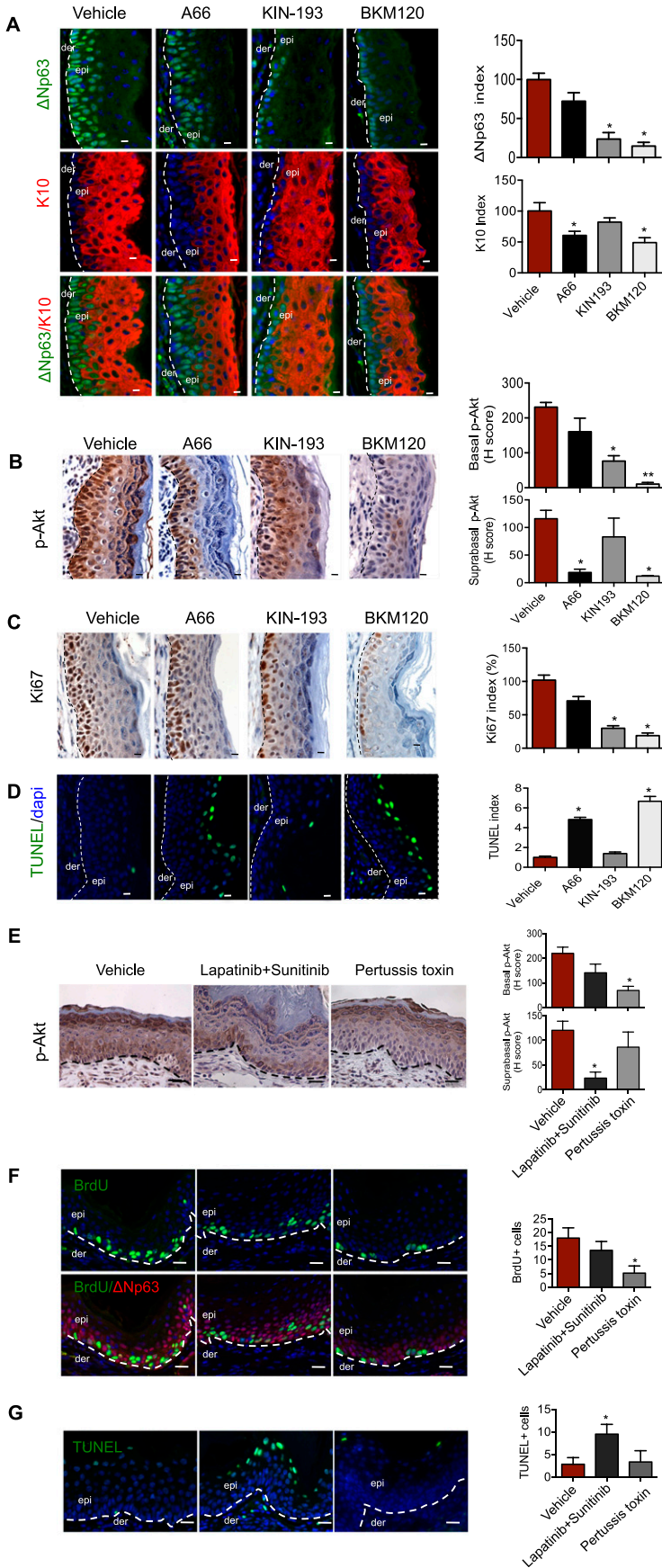


Figure 4. Effects of p110 α -selective inhibitor (A66) and p110 β -selective inhibitor (KIN-193) on PHTS of *Pten* Δ mice. (A) IF staining analyses of Δ Np63 and K10 in epidermal sections of ears from hamartoma-bearing *Pten* Δ mice 3 d after treatment with vehicle, A66 (100 mg/kg, once per day [QD]), KIN-193 (20 mg/kg, twice per day [BID]), or BKM120 (45 mg/kg, QD). Dashed lines denote the dermo/epidermal border. Bar, 20 μ m. The number of Δ Np63- and K10-positive cells in a 100- μ m length of epidermis of wild-type mice was used as a control (defined as 1). The bar graph represents the Δ Np63 or K10 index of each genotype relative to the control (\pm SEM; $n = 6$ for each genotype). (*) $P < 0.01$ (Student's t -test). (B) IHC analysis of p-Akt in epidermal sections of ears from hamartoma-bearing *Pten* Δ mice 3 d after treatment with vehicle, A66 (100 mg/kg, QD), KIN-193 (20 mg/kg, BID), or BKM120 (45 mg/kg, QD). Dashed lines mark the epidermal-dermal junction. Bar, 20 μ m. H-scores of p-Akt in basal and suprabasal epidermis are shown as mean \pm SEM ($n = 6$). (*) $P < 0.01$; (**) $P < 0.001$ (Student's t -test). (C) IHC analysis of Ki67 in the epidermal sections of ears from hamartoma-bearing *Pten* Δ mice 3 d after treatment with vehicle, A66 (100 mg/kg, QD), KIN-193 (20 mg/kg, BID), or BKM120 (45 mg/kg, QD). Bar, 10 μ m. The dashed line indicates the boundary between epidermis and dermis. The number of Ki67-positive cells in a 100- μ m length of epidermis of *Pten* Δ mice treated with vehicle was used as a control (defined as 100%). The bar graph shows the Ki67 index of each group of mice expressed as percentage of the control (\pm SEM; $n = 6$ for each genotype). (*) $P < 0.01$ (Student's t -test). (D) TUNEL assay of epidermal sections of ears from hamartoma-bearing *Pten* Δ mice 3 d after treatment with vehicle, A66 (100 mg/kg, QD), KIN-193 (20 mg/kg, BID), or BKM120 (45 mg/kg, QD). The dashed line indicates the boundary between epidermis and dermis. The number of TUNEL-positive cells in a 100- μ m length of epidermis of *Pten* Δ mice treated with vehicle was used as a control (defined as 1). The bar graph shows the TUNEL index of each group of mice expressed as percentage (\pm SEM) over the control ($n = 6$ for each genotype). Bar, 10 μ m. (*) $P < 0.001$ (Student's t -test). (E) IHC analysis of p-Akt in epidermal sections of ears from hamartoma-bearing *Pten* Δ mice 3 d after treatment with vehicle, a combination of lapatinib (100 mg/kg, QD) and sunitinib (40 mg/kg, QD), or pertussis toxin (20 μ g/kg, QD). Dashed lines mark the epidermal-dermal junction. Bar, 20 μ m. H-scores of p-Akt in basal and suprabasal epidermis are shown as mean \pm SEM ($n = 6$). (*) $P < 0.01$ (Student's t -test). (F) IF staining analyses of BrdU-positive cells and Δ Np63-positive cells in epidermal sections of ears from hamartoma-bearing *Pten* Δ mice 3 d after treatment with vehicle, a combination of lapatinib (100 mg/kg, QD) and sunitinib (40 mg/kg, QD), or pertussis toxin (20 μ g/kg, QD). Dashed lines denote the dermo/epidermal border. Bar, 20 μ m. The number of BrdU-positive cells in a 100- μ m length of epidermis of mice as indicated. (\pm SEM; $n = 6$). (*) $P < 0.01$ (Student's t -test). (G) TUNEL assay of epidermal sections of ears from hamartoma-bearing *Pten* Δ mice 3 d after treatment with vehicle, a combination of lapatinib (100 mg/kg, QD) and sunitinib (40 mg/kg, QD), or pertussis toxin (20 μ g/kg, QD). The dashed line indicates the boundary between epidermis and dermis. The number of TUNEL-positive cells in a 100- μ m length of epidermis of *Pten* Δ mice. Bar, 20 μ m. (*) $P < 0.01$ (Student's t -test).

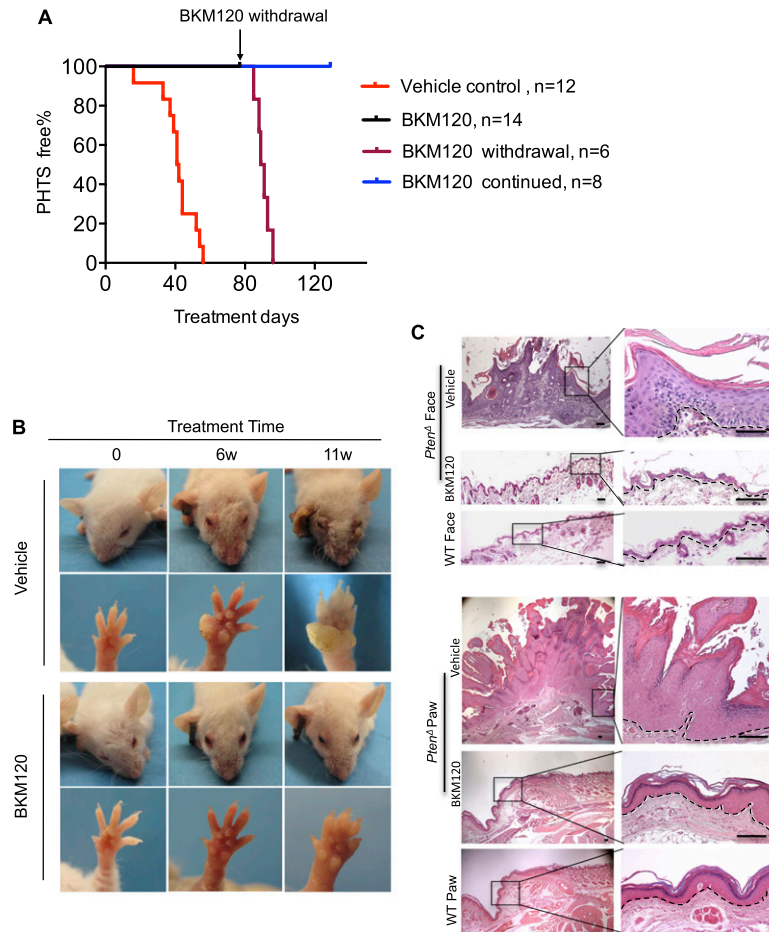


Figure 5. Early administration of BKM120 prevents the development of skin PHTS in mice. (A) KM curves for *Pten*^{-/-} mice treated with BKM120 (25 mg/kg, QD) (black line; *n* = 14) or vehicle only (red line, *n* = 12) starting at 3 wk of age. BKM120 was withdrawn from six of BKM120-treated mice at the end of the 11th week of treatment as shown by the dark-red line. BKM120 treatment was continued for the remaining eight mice for an additional 6 wk, as shown by the blue line. (B) Representative photos showing the appearance of the faces and paws of *Pten*^{-/-} mice at 6 and 11 wk post-treatment with vehicle or BKM120 (25 mg/kg, QD). Treatment was started at 3 wk of age. (C) Representative H&E staining of tissue sections of face skin and paw regions from *Pten*^{-/-} mice upon completion of 11 wk of treatment with vehicle or BKM120 (25 mg/kg, QD) beginning at 3 wk of age. Dashed lines denote the dermo/epidermal border. Bar, 100 μ m.

administered daily, and mice were monitored for morphological skin changes. We confirmed by Western blot analyses that BKM120 administered at 45 mg/kg abrogated p-Akt levels signals in *Pten*-deficient hamartoma epidermis (Supplemental Fig. 13). Our initial study was performed on a small cohort of *Pten*^{-/-} mice consisting of only two mice with advanced skin hamartomas. Prior to treatment, mouse #J845 (15 wk old) displayed severe skin lesions associated with hyperkeratosis on its face and paws (Fig. 6A). Remarkably, the facial lesions on this mouse disappeared, and the papillomas on its paw were dramatically reduced after 2 wk of BKM120 treatment; the skin conditions continued to improve up to 4 wk of treatment (Fig. 6A). Similarly encouraging results were observed on mouse #J596 (14 wk old) after treatment with this PI3K inhibitor. A large tumor that had developed on the left ear shrank dramatically after 2 wk of BKM120 treatment; after 4 wk of treatment, this tumor was completely eradicated (Fig. 6A). We conducted the same treatment regimen on a group of six *Pten*^{-/-} mice (between 12 and 16 wk of age) bearing severe PHTS on their faces, ears, and paws. Extraordinarily, BKM120 treatment led to rapid regression of all skin lesions on the faces, ears, and paws of treated mice. Significant morphological changes in these mice started to manifest 1 wk after the treatment, and their skin condition continued to improve, reaching

a near-normal state after 4 wk of treatment (Fig. 6A; Supplemental Fig. 14). Consistent with these findings, histological analyses of skin biopsies from the faces and paws of PHTS-bearing mice before and after treatment demonstrated that the drug resulted in a dramatic transformation in these mice, with skin progressing from a disordered condition to a near-normal state as assessed by histology and various molecular staining, including Δ Np63/K10, p-Akt, and Ki67 (Fig. 6B; Supplemental Fig. 15). Thus, continuously up-regulated PI3K activity appears to be essential for the maintenance of these skin hamartomas induced by *Pten* loss, and pharmacological inhibition of class I PI3K activity effectively reverses PHTS in mice.

Discussion

Our studies using genetic mouse models revealed that a balance between PI3K and *Pten* activity is critical in skin homeostasis. While *Pten* functions as a dual protein/lipid phosphatase and has also been found to have phosphatase-independent modes of action, we showed that loss of its lipid phosphatase activity in particular is largely responsible for its role in skin PHTS. The consequences of the loss of *Pten* activity can be suppressed or reversed through genetic ablation of *p110 α* and *p110 β*

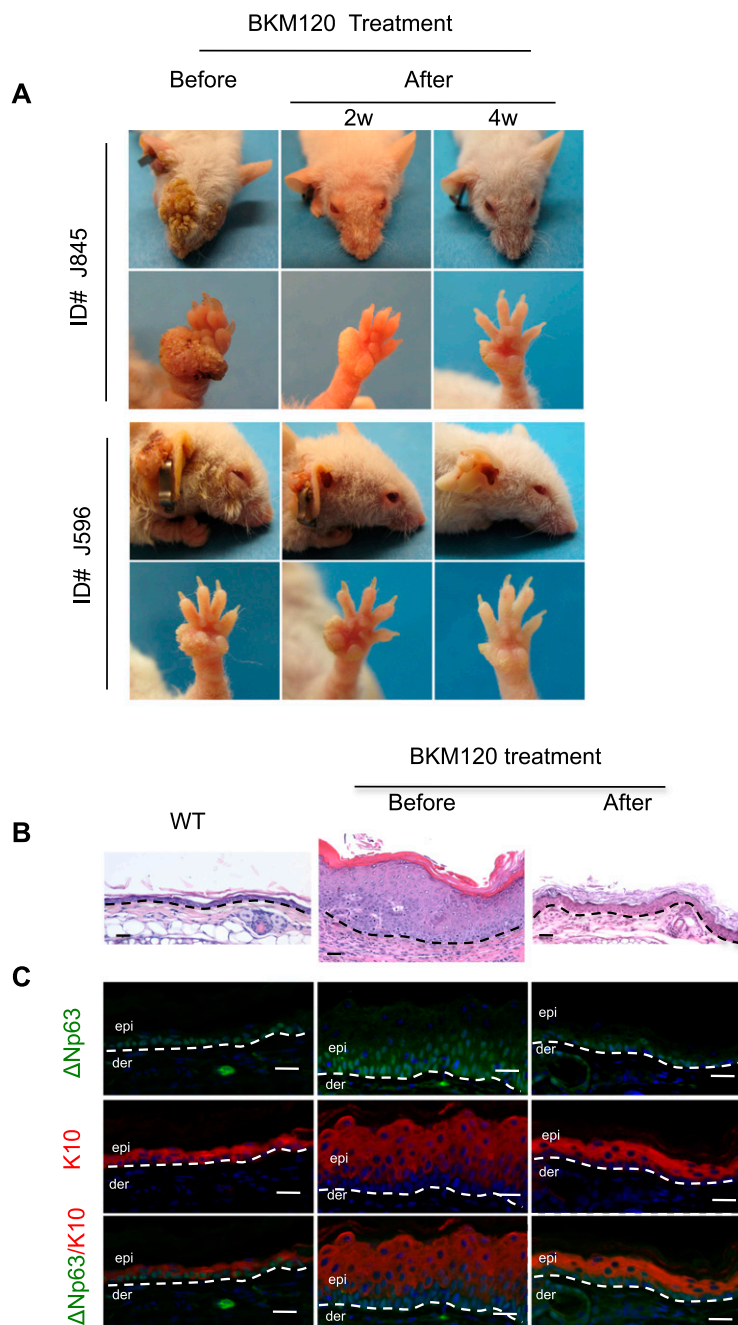


Figure 6. Administration of BKM120 relieves PHTS in *Pten*^Δ mice. Hamartoma-bearing *Pten*^Δ mice were treated with vehicle or BKM120 (45 mg/kg, QD) for 4 wk. (A) Representative images of two *Pten*^Δ mice (ID #J845 and #J596) before and after 2 wk and 4 wk of treatment with BKM120. (B) Representative H&E staining of tissue sections of ears from hamartoma-bearing *Pten*^Δ mice before and after 4 wk of treatment with BKM120 (45 mg/kg, QD). Dashed lines denote the dermo/epidermal border. Bar, 20 μ m. (C) IF staining analyses of Δ Np63 and K10 in epidermal sections of ears from hamartoma-bearing *Pten*^Δ mice treated with vehicle or BKM120 (45 mg/kg, QD) for 4 wk. Dashed lines denote the dermo/epidermal border. Bar, 20 μ m.

isoforms of class Ia PI3K or by their pharmacological inhibition via a pan-class I PI3K inhibitor. The more nuanced roles of the two commonly expressed PI3K isoforms in skin contrasts with our previous findings that ablation of *p110 β* was sufficient to block *Pten* loss-induced PIN in the anterior lobes of mouse prostates (Jia et al. 2008).

Here we report an interesting and surprising “division of labor” in a spatially defined manner between the two isoforms in maintaining skin epithelium, as depicted in the diagram in Figure 7A. The unique cellular architecture of the *Pten*-null epidermis, dramatically thickened skin, and increased epidermal layers, provide us with an

exceptional platform to determine the role that p110 α or p110 β activity plays in PHTS by examining the consequences of their inhibition either genetically or pharmacologically (Fig. 7B). In the *Pten*-null epidermis where p110 α is also genetically or pharmacologically inactivated, p110 β becomes the major PI3K player that is unrestrained by its lipid phosphatase, resulting in a hyperproliferative epithelium featuring an expansion of its basal cell layers (Fig. 7B, third panel). In contrast, when *Pten* and p110 β are absent in the epidermis, uncontrolled p110 α activity sustains survival of terminally differentiated cells such that these cells fail to undergo apoptosis

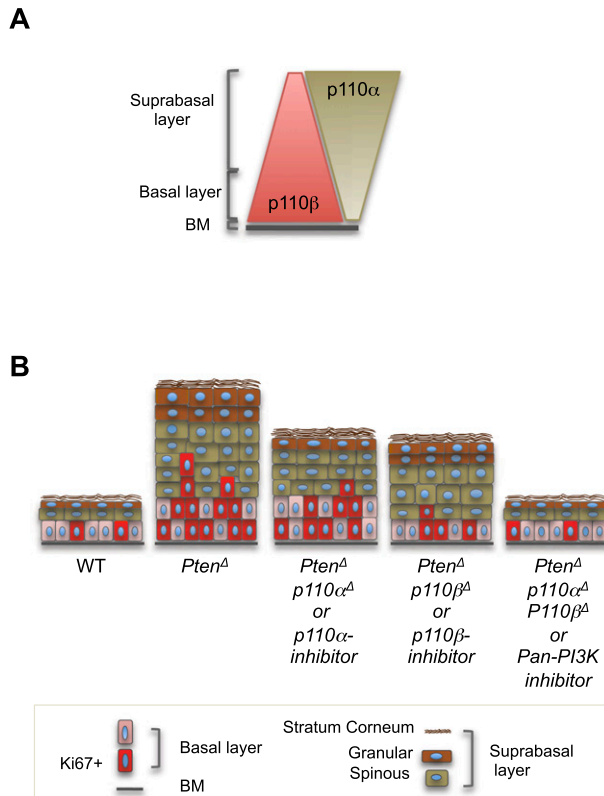


Figure 7. (A) Summary of spatially differential roles of p110 α and p110 β isoforms in skin epidermis. (B) Diagram representation of skin histology in mice and the extent of p110 α 's and p110 β 's role in the epidermis lacking *Pten*.

as they reach the skin surface, resulting in the accumulation of terminally differentiated suprabasal cells (Fig. 7B, fourth panel). The PHTS skin can only be restored to normal by suppressing both PI3K isoforms. Thus, *Pten* has an essential role in counterbalancing the kinase activities of both p110 isoforms to maintain skin homeostasis and prevent the development of PHTS.

While our finding of the spatially differential expression patterns of p110 α and p110 β provides a mechanistic understanding of the specific roles of these two isoforms in epidermis, clearly, additional factors are also at play in the regulation of these PI3K isoforms in the skin epidermis. Since *Pten* loss simply removes a "brake" from the PI3K pathway, the specific roles of p110 isoforms in various pathogenic states induced by *Pten* loss can be influenced by tissue microenvironment cues that mediate the activation of PI3K; e.g., growth factors, chemokines, extracellular matrix proteins, and other input signals. Indeed, we found that broad-spectrum inhibition of RTKs resulted in reduced p-Akt in suprabasal cells and increased apoptosis of granular cells in *Pten*-null epidermis. Conversely, a general GPCR signaling inhibitor (pertussis toxin) largely reduced Akt phosphorylation and cell proliferation in basal cells of *Pten*-null epidermis. Thus, the p110 α -dependent signal is likely due to activation of an RTK in the suprabasal layers, while the p110 β -dependent signaling likely arises

from either GPCR or integrin-mediated signaling or both. Notably, in the skin, only basal cells contact the basement membrane and hence feature integrin engagement. However, precisely how diverse signals activate certain PI3K isoforms and how these activities are coordinated with each other and with *Pten* to control this pathway in different tissues and cell types remain to be determined.

Importantly, our work holds potential for translating insights from mouse genetics into human therapy. Emerging from our study is the potential benefit of pharmacological PI3K inhibitors for PHTS patients as both therapeutic agents and chemopreventive agents (at reduced doses) with meaningful clinical outcomes. The newly identified spatially divergent roles for p110 α and p110 β in skin suggest that the use of PI3K inhibitors to treat these skin lesions can be versatile. For example, if long-term systemic use of a pan-inhibitor proved to have adverse metabolic effects even at the low doses that we found beneficial in mice, one might take a combination approach using topical application of a p110 α inhibitor with the systemic administration of a drug directed against p110 β . Since p110 α is essential for the metabolic functions of insulin and for glucose homeostasis (Sopasakis et al. 2010), external application may provide safer and more effective therapy for PHTS patients.

Materials and methods

Mice

K14Cre (Jonkers et al. 2001), *Pten*^{L/L} (Lesche et al. 2002), *p110α*^{L/L} (Zhao et al. 2006), and *p110β*^{L/L} (Jia et al. 2008) mice were backcrossed 10 generations to the FVB/N background before being intercrossed to generate experimental mice for this study. All animals were housed and treated in accordance with protocols approved by the Institutional Animal Care and Use Committees at Dana-Farber Cancer Institute and Harvard Medical School.

PHTS severity score

The progression of mucocutaneous lesions in mice was monitored according to the operational diagnosis criteria of PHTS (Nelen et al. 1996; Eng 2000; Laury et al. 2011). The facial papillomatous, acral keratoses, and palmoplantar keratoses (limbs and paws) were each graded from 0 (normal) to 4 (severe), and a summed score was calculated accordingly. Each grade was assigned by histological assessment. The skin samples, which were graded >2 in each area, were diagnosed as papilloma, a benign skin hamartoma. Papillomas in more than two areas in the body, or a summed score of 4, were considered PHTS.

Histology, IHC, and immunofluorescence (IF)

For histology analyses, formalin-fixed tissue sections were embedded in paraffin, sectioned, and stained with hematoxylin and eosin by the Dana Farber/Harvard Cancer Center Rodent Histopathology Core. Antibodies used for IHC and IF were Ki67 (Vector Laboratories), p-Akt (Cell Signaling), p63 (Santa Cruz Biotechnology), Δ Np63 (Biolegend), loricrin (Abcam), K5 (Convance), K10 (Convance), and rat anti-BrdU (AbD seroTec). For BrdU analysis, mice were injected with 50 mg BrdU per kilogram

of body weight 2 h prior to sacrifice. For IF staining, after incubation with the primary antibody, the secondary antibodies conjugated to different Alexa-Fluor dyes (488 and 594) were applied at a dilution of 1:1000 in PBS. For nuclear counterstaining, DAPI was used. TUNEL assays were performed with the In Situ Cell Death Detection kit (Roche) according to the manufacturer's instructions. Samples were then counterstained with DAPI and analyzed under a fluorescent microscope. For quantification of IHC and IF, six images were taken per skin section, and the positive cells were counted.

Quantification of p-Akt

Quantification of p-Akt in IHC analyses was performed using CellVigene (VigeneTech, Inc.) protocols. The RGB colors were deconvoluted to haem (blue) and dab (brown). The haem component was used to identify nuclei. The dab component was used to quantify nuclear, cytoplasmic, and total p-Akt in basal and suprabasal regions. Background subtraction was applied to each image to minimize overstaining effects. The background values were determined by drawing a region outside the basal and suprabasal areas that CellVigene software uses to autodetermine the background value in the region. The basal and suprabasal regions were classified using supervised machine learning. Typical four-level histogram scoring and mean nuclear and cytoplasmic intensities were used to calculate the percentage of a score level. A histogram score (H-score), which was equal to $1 \times (1+) \% + 2 \times (2+) \% + 3 \times (3+) \%$, ranging from 0 to 300, was used to measure p-Akt.

Western blotting

Primary keratinocytes were isolated as described (Dlugosz et al. 1995). Briefly, nonhairy ear skin from mice was washed in betadine and 70% ethanol. The two skin flaps of each ear were separated carefully and trypsinized for 1.5 h by floating on 1% trypsin at 37°C. The epidermis was scraped off the dermis. Cells of epidermis were then lysed in 1% Nonidet P-40 lysis buffer supplemented with protease and phosphatase inhibitors (Roche). Equal amounts of proteins were resolved on SDS-PAGE gels and transferred to nitrocellulose membranes for Western blot analysis. The primary antibodies used were p-Akt (Ser473), p-Akt (Thr308), Akt, p110 α , p110 β , and Pten (all from Cell Signaling Technology). Immunofluorescently labeled anti-mouse IgG (Rockland Immunochemicals) and anti-rabbit IgG (Molecular Probes) were used to visualize Western blots on an Odyssey scanner (Li-Cor). Quantification of band intensities was performed using Odyssey 3.0 software.

FACS analysis

Keratinocytes were isolated from the ear epidermis of 8-wk-old wild-type or *Pten*^{-/-} mice. Single-cell suspensions in PBS containing 2% FCS were incubated with primary antibodies for 30 min on ice prior to FACS analysis. Primary antibodies used for FACS analysis were APC α 6-integrin antibody (R&D Systems) and PE anti-mouse CXCR3 (Biolegend). Cell sorting was performed with BD FACSAria IIu and analyzed with FACSDiva software or FlowJo.

Quantitative RT-PCR

Total RNAs were extracted from epidermal cells using TRIzol reagent and used to synthesize cDNA with SuperScript III first strand synthesis kit (Invitrogen). Quantitative PCR reactions were performed with SYBR Green PCR Master Mix (Applied

Biosystems). Primer sequences of *p110 α* (NM_008839) were as follows: forward, 5'-TTCTCTGGAAATGCAGACC-3'; and reverse, 5'-GTGGACAGCATCCCTGTAAC-3'. Primers for *p110 β* were purchased from SABioscience (PPM05089A). The levels of amplified transcripts were normalized to that of control transcript 18s rRNA (Bouras et al. 2008).

Inhibitor administration

BKM120 (Novartis Pharmaceuticals) was reconstituted in 1 vol of NMP (1-methyl-2 pyrrolidone; Sigma) and 9 vol of PEG-300 (polyethylene glycol 300; Fluka Analytical) (Maira et al. 2012). Mice were dosed with this compound formulation at 25 or 45 mg/kg once per day (QD) by oral gavage. A66 was administered in 10% 2-hydroxypropyl- β -cyclodextrin (Sigma-Aldrich) in water as described (Jamieson et al. 2011). Mice were dosed with 100 mg/kg QD by intraperitoneal (IP) injection. KIN-193 was dissolved in 7.5% NMP and 40% PEG400 (polyethylene glycol 300; Fluka Analytical) as described (Ni et al. 2012). Mice were dosed with 20 mg/kg twice per day (BID) by IP injection. Lapatinib was dissolved in 0.5% hydroxypropyl methylcellulose and 1% Tween80 as described (Konecny et al. 2006). Sunitinib (Sigma Aldrich) was prepared in dextrose water (Zhang et al. 2009). Mice were gavaged with lapatinib (100 mg/kg) and sunitinib (40 mg/kg) QD (Sun et al. 2011). Pertussis toxin (Sigma Aldrich) was dissolved in PBS. Mice were dosed with 20 μ g/kg QD by IP injection as described (Chen et al. 2007).

Statistics

Comparison of the Kaplan-Meier analyses was performed with the log-rank test. For all bar graphs, the Student's *t*-test was used to determine a *P*-value, and data are presented as mean \pm SEM.

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