ΔNp63 α regulates keratinocyte proliferation by controlling PTEN expression and localization

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ΔNp63 α , implicated as an oncogene, is upregulated by activated Akt, part of a well-known cell survival pathway. Inhibition of Akt activation by phosphatase and tensin homolog deleted on chromosome 10 (PTEN) and the presence of putative p63-binding sites in the *pten* promoter led us to investigate whether ΔNp63 α regulates PTEN expression. Knockdown of ΔNp63 α led to increases in PTEN levels and loss of activated Akt, while overexpression of ΔNp63 α decreased PTEN levels and elevated active Akt. The repression of PTEN by ΔNp63 α occurs independently of p53 status, as loss of ΔNp63 α increases PTEN expression in cell lines with and without functional p53. In addition, decreased levels of ΔNp63 α resulted in an increase in nuclear PTEN. Conversely, *in vivo* nuclear PTEN was absent in the proliferative basal layer of the epidermis where ΔNp63 α expression is highest. Additionally, we show that in keratinocytes a balance between ΔNp63 α and PTEN regulates Akt activation and maintains normal proliferation rates. This balance is disrupted in non-melanoma skin cancers through increased ΔNp63 α levels, and could enhance proliferation and subsequent neoplastic development. Our studies show that ΔNp63 α negatively regulates PTEN, thereby providing a feedback loop between PTEN, Akt and ΔNp63 α , which has an integral role in skin cancer development. *Cell Death and Differentiation* (2011) 18, 1924–1933; doi:10.1038/cdd.2011.73; published online 3 June 2011

The p53 transcription factor family consists of the tumor suppressor p53 and the homologous p63 and p73. Unlike p53, p63 is essential for normal epidermal stratification and the proliferative potential of the epithelial stem cells.^{1,2} p63 exists as various isoforms with contrasting functions.² The TA isoforms (TAp63 α , TAp63 β and TAp63 γ) have a full-length N-terminal transactivation domain, whereas the ΔN isoforms (ΔNp63 α , ΔNp63 β and ΔNp63 γ) have a short but distinct transactivation domain. All isoforms have a DNA-binding domain that shares high homology with p53 that allows p63 proteins to bind to p53 DNA-binding sites.^{3,4}

Similar to p53, the TA isoforms of p63 and p73 can promote apoptosis and growth arrest through the induction of anti-proliferative genes. In contrast, the ΔN isoforms have been shown to induce pro-survival genes and inhibit anti-proliferative genes.^{4–6} Several studies indicate that ΔNp63 α , the predominant isoform in adult tissue, may function as an oncogene as it can exert a dominant-negative effect over p53 and the TAp63 and TAp73 isoforms.² Additionally, ΔNp63 α is frequently overexpressed in a variety of squamous cell (SCC) and basal cell carcinomas (BCCs).^{7,8}

The survival factor Akt can increase ΔNp63 α levels and in turn, ΔNp63 α protects against UV-B-induced apoptosis via Akt activation.^{9,10} However, the mechanism behind the positive feedback loop between ΔNp63 α and Akt has not been described. Akt activation can be negated by phosphatase and tensin homolog deleted on chromosome 10 (PTEN). PTEN

dephosphorylates phosphatidylinositol 3,4,5-trisphosphate, thereby inhibiting the signaling cascade that activates Akt. Additionally, PTEN can undergo nuclear localization to induce G1 arrest independently of Akt.^{11–13} The *pten* gene is induced by p53 and, considering that ΔNp63 α can oppose p53 function and bind to p53 responsive elements (REs), it is likely that ΔNp63 α may negatively regulate PTEN.^{2,4,14} In this study we report that loss of ΔNp63 α results in upregulation of PTEN and subsequent decrease in Akt activation and cell proliferation. We demonstrated that a balance between ΔNp63 α and PTEN expression is required to maintain the rate of normal proliferation, and disruption of this balance potentiates cell growth via activation of the Akt pathway. Finally, we showed that the ratio of ΔNp63 α to PTEN expression was significantly perturbed in human non-melanoma skin cancers.

Results

ΔNp63 α negatively regulates PTEN expression. To address the potential functional interaction between endogenous p63 and PTEN, we first characterized the predominant p63 isoform in A431 and HaCaT. HaCaT cells were chosen because they are non-tumorigenic and represent an intermediary between normal and cancerous keratinocytes. A431 cells were derived from an atypical SCC of the vulva epidermis. Consistent with earlier reports, we

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Abbreviations: ΔNp63 α , deltaNp63alpha; PTEN, phosphatase and tensin homolog deleted on chromosome 10; HEKs, primary human epidermal keratinocytes; RE, response element; CA-Akt, constitutively active Akt; BCC, basal cell carcinoma; SCC, squamous cell carcinoma; MFI, mean fluorescence intensity

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demonstrated that Δ Np63 α is the predominant isoform in both HaCaT and A431 cell lines.^{15,16} A faint band corresponding to Tap63 γ was detected upon immunoblot analysis using a TA-specific antibody (Supplementary Figure S1). To determine the effect of Δ Np63 α on PTEN, we knocked down p63 in both mutant p53 (A431 and HaCaT, Figure 1a) and wild-type p53 (primary human keratinocytes, Figure 1b) backgrounds. Silencing p63 with multiple pan-p63 siRNAs led to increased PTEN levels independently of p53 status (Figures 1a and b). Increased IGFBP-3 levels, known to be negatively regulated by Δ Np63 α , served as a positive control.⁵ We confirmed that the pan-p63 siRNAs led to knockdown of both Tap63 and Δ Np63 by isoform-specific qRT-PCR (Supplementary Figure S1C). Loss of only Δ Np63 α by a Δ Np63-specific siRNA still increased PTEN expression, confirming that Δ Np63 α is the main regulator of PTEN (Figure 1c). Although the increase in PTEN protein seems modest, minute changes in PTEN protein levels have profound biological impact and thus the modest increase we observe is significant.¹⁷ PTEN levels were also elevated

in E1A-transformed murine embryonic fibroblasts (MEFs) and primary MEFs obtained from p63 $^{-/-}$ mice when compared with wild-type mice (Figure 1d). A minor amount of residual p63 mRNA was detectable in the p63 $^{-/-}$ MEFs because of the location of primers used for qRT-PCR, which produces a non-functional protein.¹⁸ Taken together, the results from these sets of experiments indicate that PTEN is negatively regulated by Δ Np63 α .

Δ Np63 α promotes survival of squamous carcinomas of the head and neck by repressing p73-dependent pro-apoptotic effects.¹⁹ The levels of both TA and Δ Np73 (Supplementary Figure S2) were examined, and Δ Np73 was the predominant isoform observed, suggesting that these cells may have a deficient p73-dependent pro-apoptotic program (Supplementary Figure S2). However, low levels of Tap73 α and β were still detectable. To rule out the possibility that silencing Δ Np63 α was inducing PTEN expression by reversing its dominant-negative effects on Tap73 or Tap63, we transfected cells with siRNA to Δ Np63 α and/or siRNA specific to either Tap63 or Tap73 (Supplementary Figure S3). Importantly, we did not

Figure 1 Δ Np63 α negatively regulates PTEN expression. (a) A431, HaCaT and (b) HEK cells were transfected with non-silencing control (NSC) siRNA or two to three different siRNAs targeting p63. Total RNA was extracted and transcript levels of p63, PTEN and IGFBP-3 were analyzed by qRT-PCR. Y-axis represents the fold change in PTEN and p63 transcript levels relative to NSC-transfected cells. Immunoblots of p63 and PTEN in A431 and HaCaT cells transfected as in (a) and HEKs as in (b) are shown in the bottom panels. Fold change in protein relative to NSC is listed above each band. (c) HaCaT cells were transfected with NSC or siRNA specific to Δ Np63 isoforms of p63. Total RNA and protein were extracted and analyzed as in (a). (d) Total RNA extracted from both primary and E1A-transformed MEFs obtained from wild-type and p63 $^{-/-}$ mice were subjected to qRT-PCR to detect transcript levels in the top panel. The bottom panel represents immunoblot analysis of whole-cell extracts from wild-type and p63 $^{-/-}$ MEFs. Note, p63 was undetectable at the protein level and near the limit of detection for qRT-PCR in p63 $^{-/-}$ MEFs. Error bars represent standard deviation. * $P < 0.05$ compared with control cells in all panels

observe a reduction in PTEN expression upon knockdown of either the Tap63 or the Tap73 isoforms, ruling out the possibility that the TA isoforms in these cells were positively transcriptionally regulating PTEN (Supplementary Figure S3). Dual knockdown of Δ Np63 α and Tap63 or Δ Np63 and Tap73 led to significant PTEN inductions similar to silencing Δ Np63 alone, concluding that PTEN induction observed upon silencing Δ Np63 α occurs independently of any effects on Tap63 or Tap73 activity.

PTEN is a direct target of Δ Np63 α . Examination of the *pten* promoter revealed the presence of several putative p63 REs in addition to the p53 RE described previously (Figure 2a).¹⁴ To define the region within the *pten* promoter to which Δ Np63 α binds, we carried out chromatin immunoprecipitation (ChIP) assays in A431 and HaCaT cells using antibodies that recognize all p63 isoforms or only the p63 α isoforms. p63 bound to the same locations of the *pten* promoter with similar affinities in both cell lines (Figure 2b). Similar to other genes, p63 also bound the p53 RE (region C, Figure 2b).⁴ Region D contained two p63-specific REs, yet failed to demonstrate binding. Region E only showed binding when immunoprecipitated with the pan-p63 antibody, indicating weak binding to this RE. Because mutant p53 has been demonstrated to complex with p63 and affect its binding to transcriptional targets, we also evaluated the binding of p53 to the *pten* promoter.^{20,21} p53 bound nearly all of the same regions of the *pten* promoter as p63 (Supplementary Figure S4). Consistent with previous reports, the mutant p53 in

A431 cells failed to bind the p21 promoter, suggesting that this mutation disrupts the normal binding pattern of p53.²² Contrary to previous studies, our results suggest that although mutant p53 may interact with p63, it does not prevent p63 from binding to the promoter of *pten*.^{20,21} These results show that endogenous Δ Np63 α is capable of binding to the *pten* promoter in multiple locations unique to p63 in addition to binding the p53 RE.

To define the minimal region within the *pten* promoter critical for Δ Np63 α -mediated repression of PTEN, we analyzed the effect of Δ Np63 α on a luciferase reporter containing the full-length *pten* promoter (PTEN-Luc). p63 and p53 null H1299 cells transfected with Δ Np63 α led to a statistically significant reduction of PTEN-Luc reporter activity when compared with control cells (Figure 2c, top panel). Silencing Δ Np63 α in HaCaT cells led to a modest but reproducible increase in the PTEN-Luc reporter activity (Figure 2c). We next generated minimal luciferase reporters containing only the putative p63 REs found in the regions within the *pten* promoter that showed p63 binding by ChIP (Figure 2b). Co-transfection of these reporters with Δ Np63 α exhibited a significant reduction in reporter activity only with the minimal reporter corresponding to region B (Figure 2c, lower panel). These results suggest that negative regulation of PTEN by Δ Np63 α is likely due to binding of Δ Np63 α to region B of the *pten* promoter.

A balance between Δ Np63 α and PTEN controls cell proliferation. To examine whether Δ Np63 α promotes cell

Figure 2 PTEN is a direct target of Δ Np63 α . (a) Schematic of putative p63-binding sites and documented p53-binding sites in the *pten* promoter (accession no. AF067844). (b) ChIP assay was performed on A431 and HaCaT cells, wherein chromatin was immunoprecipitated with normal IgG control antibody or antibodies that recognize all p63 isoforms or p63 α isoforms as indicated. Eluted DNA was PCR-amplified with primers specific for multiple regions of the *pten* promoter as shown in (a). The p21 promoter was used as a positive control for p63 binding. (c) The top left panel shows dual luciferase assays performed on H1299 cells co-transfected with the full-length PTEN-Luc reporter and empty vector or Δ Np63 α . The top right panel shows dual luciferase assays performed on HaCaT cells after knockdown of p63 by siRNA. Y-axis represents fold change in relative luciferase units compared with control-transfected cells. Representative immunoblots are shown below as proof of adequate overexpression or knockdown. The bottom panel shows dual luciferase assays in H1299 after co-transfection of the designated minimal PTEN-luc reporter and empty vector or Δ Np63 α . The sequence corresponding to region B is shown below, with the core sequence of the putative p63 REs bolded and underlined. Lined up underneath each p63 half site is the canonical p63 RE, where R denotes purines; W, A or T; and Y, pyrimidines. Error bars represent standard deviation. * $P < 0.05$ compared with control cells

proliferation by inhibiting PTEN, colony-formation assays were carried out after silencing Δ Np63 α and/or PTEN. Silencing Δ Np63 α reduced colony formation, whereas silencing PTEN led to an increase in total colony number and in colony size. Interestingly, no statistical difference in colony number was observed when both Δ Np63 α and PTEN were silenced when compared with control-transfected cells (Figure 3a). Silencing of Δ Np63 α and PTEN was confirmed by immunoblot analysis as shown in Figure 3a. Consistent with this, cell proliferation determined by MTS assay, which measures the amount of actively metabolizing cells (Figure 3b), or cell viability determined by trypan blue

exclusion (Figure 3c) also demonstrated that silencing both Δ Np63 α and PTEN resulted in cell growth rates comparable to control cells. These results, based on three independent measures of cell growth, suggest that a critical balance between PTEN and Δ Np63 α must be reached to maintain normal proliferation as perturbation of either PTEN or Δ Np63 α levels affects proliferation.

Activation of the Akt pathway via Δ Np63 α -mediated PTEN repression. We next examined whether Δ Np63 α -mediated regulation of PTEN affects Akt activation. Increasing levels of Δ Np63 α led to a significant,

Figure 3 A balance between Δ Np63 α and PTEN controls cell proliferation. (a) A431 and HaCaT cells were transfected with NSC siRNA, p63 siRNA and/or PTEN siRNA as indicated. The left panel shows representative images of crystal violet-stained colonies at 3 days post re-seeding of cells transfected with designated siRNA or NSC. y-axis represents total number of colonies from (a) as counted with ImagePro 6.2. Error bars represent standard deviation. * $P < 0.05$. Adequate knockdown was confirmed by immunoblot analysis and a representative experiment is shown. (b and c) A431 and HaCaT cells were transfected with NSC siRNA, p63 siRNA and/or PTEN siRNA as indicated and replated 24 h post transfection. Cells were subjected to MTS assays (b) or trypan blue staining (c) on the indicated day. y-axis represents total cell number (c). Error bars representing standard deviation within a single representative experiment

dose-dependent reduction in endogenous PTEN protein levels with a concomitant increase in phosphorylated Akt (p-Akt) levels (Figure 4a). Induction of p-Akt by Δ Np63 α is dependent on PTEN as overexpression of Δ Np63 α in PTEN null PC3 cells failed to increase p-Akt levels (Figure 4b). We next wanted to determine whether Δ Np63 α and constitutively active Akt (CA-Akt) had a synergistic effect on reduction of PTEN levels.¹⁰ Δ Np63 α alone led to a dramatic reduction in PTEN expression, while CA-Akt alone led to a modest decrease in PTEN (Figure 4c). Co-expression of Δ Np63 α and CA-Akt led to an enhancement of Δ Np63 α , and PTEN was further reduced as compared with CA-Akt alone (Figure 4c). The ability of Δ Np63 α alone to reduce PTEN expression suggests that Δ Np63 α amplification is comparable to mutations in the PI3K/Akt pathway with regard to deregulating tumor suppression. Silencing Δ Np63 α led to an increase in endogenous PTEN corresponding to decreased p-Akt levels (Figure 4d). Consistent with the return to normal growth rates when both Δ Np63 α and PTEN were silenced in Figure 3, loss of both Δ Np63 α and PTEN results in p-Akt levels similar to those of control-treated cells (Figure 4e). These data demonstrate that Δ Np63 α and PTEN work in concert to modulate p-Akt levels to control proliferation rates.

Δ Np63 α suppresses nuclear PTEN. Although cytoplasmic PTEN antagonizes the Akt pathway, nuclear PTEN is associated with cell-cycle arrest that may be independent of Akt.^{13,23} Therefore, we examined whether silencing Δ Np63 α affects both nuclear and cytoplasmic PTEN levels.

A significant increase in nuclear PTEN was observed when Δ Np63 α was silenced (Figure 5a). Similarly, silencing Δ Np63 α also led to increased nuclear PTEN at a cellular level, and non-silenced cells showed little nuclear PTEN (Supplementary Figure S5). Although there were increases in nuclear and cytoplasmic PTEN levels with a corresponding decrease in p-Akt levels, only the decrease in cytoplasmic p-Akt was statistically significant, supporting the hypothesis that nuclear PTEN has some functions that are independent of Akt (Figure 5a). Δ Np63 α is often amplified in SCCs, so we next explored the effect of Δ Np63 α overexpression on nuclear PTEN. As seen in Figure 5b, ectopic Δ Np63 α led to a marked reduction of cytoplasmic PTEN with a corresponding increase in cytoplasmic p-Akt levels. A small, but reproducible, reduction in nuclear PTEN was also observed that led to a robust increase in nuclear p-Akt.

As nuclear PTEN is abundant in differentiated cells while Δ Np63 α is associated with proliferating cells, we monitored the localization of PTEN and Δ Np63 α in normal adult human and embryonic mouse epidermal tissues.¹¹ Keratinocytes of the basal layer and hair follicles, with high p63 expression, showed nearly complete nuclear exclusion of PTEN (Figure 5c). PTEN staining was only observed in the nucleus of supra-basal layer cells that no longer expressed Δ Np63 α (Figure 5c). Δ Np63 α expression is normally downregulated as keratinocytes differentiate; however, in keratinocytes of the supra-basal layer that have yet to degrade Δ Np63 α , no nuclear PTEN is observed (Figure 5c, marked by asterisk). The localization patterns of Δ Np63 α and PTEN in intact normal human and mice skin, as well as in transformed

Figure 4 Δ Np63 α affects Akt activation by regulating PTEN expression. (a) H1299 cells were transfected with increasing concentrations (0–3 μ g) of Δ Np63 α as indicated. (b) PTEN null PC3 cells were transfected with empty vector or Δ Np63 α and subjected to immunoblot analysis for the indicated proteins. (c) H1299 cells were transfected with Δ Np63 α alone or together with constitutively active Akt (CA-Akt), and whole-cell lysates were subjected to immunoblot analysis as indicated. (d) HaCaT cells were transfected with three different siRNAs targeting p63 or NSC as indicated, and whole-cell extracts were subjected to immunoblot analysis as indicated. (e) HaCaT cells were treated with NSC or siRNA against p63 and/or PTEN, and whole-cell lysates were subjected to immunoblot analysis as indicated. Fold change in protein level, normalized β -actin or to Total Akt and β -actin, is listed above each band

Figure 5 Δ Np63 α suppresses nuclear PTEN. (a) HaCaT cells were transfected with NSC or p63 specific siRNA and subjected to subcellular fractionation. Immunoblot analysis of the indicated proteins is shown in the left panel. Nuclear protein levels were normalized to the intensity of PARP, while α -tubulin served as a loading control for normalization of cytoplasmic proteins. Fold change in protein level is listed above each band. (b) H1299 cells were transfected with empty vector or Δ Np63 α before subcellular fraction and immunoblot analysis as in (a). (c) Indirect immunofluorescence was performed on normal adult human skin and E18.5 mouse skin. Representative images for nuclei (DAPI), p63 and PTEN are shown for each skin type. Basal layer cells that lack nuclear PTEN are shown by arrows, while an asterisk shows supra-basal layer cells lacking nuclear PTEN

keratinocyte cell lines, suggest that Δ Np63 α inhibits localization of PTEN to the nucleus.

Δ Np63 α and PTEN localization and expression in non-melanoma skin cancers. Elevated levels of Δ Np63 α have been reported in a variety of SCC and BCC.^{15,24,25} On the basis of the observations in Figure 5 that overexpression of Δ Np63 α could repress nuclear PTEN, we next sought to determine whether the amplification of Δ Np63 α seen in human non-melanoma cancers coincided with reduced nuclear PTEN. The expression and localization of p63 and PTEN was examined in normal skin cores ($n=9$), BCC cores ($n=11$) and SCC cores ($n=11$) using a tissue microarray (Figure 6). In normal tissue, PTEN staining was most abundant in the cytoplasm of epidermal keratinocytes (Figure 6, top left panel). A few cells with intense PTEN staining were seen in the dermis and in melanocytes, consistent with earlier reports.²⁶ It is important to remember that these cells are of mesenchymal origin and do not express any p63. In both BCC and SCC, very little

nuclear PTEN was observed in the cancerous tissue. Intense staining for PTEN within the dermis was observed in areas of solar elastosis, partially due to some non-specific uptake of secondary antibodies (Figure 6, identified as SE in the bottom panel). The levels of p63 staining in BCC and SCC were considerably higher than in normal tissue, with all cancerous islands staining positive for p63. On the basis of the results shown in Figure 5, the high levels of p63 seen in Figure 6 could possibly be contributing to the lack of nuclear PTEN in BCC and SCC.

We next sought to determine whether there was any correlation between p63 and PTEN expression in the healthy and cancerous skin samples. An increase in p63 staining, as measured by the mean fluorescence intensity (MFI), was observed in BCC and SCC when compared with normal epidermal tissue (Figure 7a). A significant correlation was observed between the MFI values of p63 and PTEN in normal tissue ($r=0.693$, $P=0.042$), and in BCC ($r=0.6687$, $P=0.00268$) and SCC ($r=0.7356$, $P=0.012$). Although there was no overall decrease in PTEN staining accompanying

Figure 6 Elevated levels of Δ Np63 α and reduced nuclear PTEN expression in basal and SCC. Indirect immunofluorescence was performed on a tissue microarray containing multiple cores of (a) normal human skin, (b) BCC and (c) SCC. Representative images of nuclei (DAPI), p63 and PTEN staining within each type of skin are shown. Areas of increased magnification to show PTEN localization are shown surrounded by a box. The regions within the tissue types are defined in the bottom panel. E, normal epidermis; D, dermis; SE, solar elastosis; BCC, basal cell carcinoma; SCC, squamous cell carcinoma

the increase in p63 in cancerous samples, there was a major perturbation in the ratio of p63 to PTEN. A statistically significant difference (ANOVA $F_{2,29} = 12.915$, $P < 0.0005$) in the ratio between p63 and PTEN MFIs was observed when normal tissue was compared with BCC and SCC (Figure 7b), highlighting the importance of the balance between Δ Np63 α and PTEN in controlling proliferation in the context of skin cancer.

Discussion

p63 is critical for epidermal stratification and the proliferative potential of the epithelial stem cells, but is also frequently overexpressed in SCCs.^{1,2,15,24,25} Skin-specific knockout of PTEN in mice leads to hyperproliferative and thickened skin.²⁷ The opposing phenotypes of PTEN and p63 in epidermal regulation suggested that these two proteins may integrate

Figure 7 Elevated levels of Δ Np63 α in basal and SCC correlate with PTEN expression. (a) Quantification of the MFI for p63 and PTEN staining, in arbitrary units, from normal epidermal tissue, BCC and SCC. Error bars represent standard error of mean. (b) The average ratio of p63 fluorescence intensity to PTEN fluorescence intensity is higher in BCC and SCC as compared with normal skin samples. Error bars represent standard error of mean. * $P < 0.0005$

into a common pathway. Δ Np63 α controls the rate of proliferation in epithelial basal cells, thereby maintaining a crucial number of basal stem cells, whereas deletion of PTEN in the basal cells leads to uncontrolled cell proliferation.^{1,28} This proliferation is considered to be an important event for initiation and progression of prostate cancer and may occur in other cancers, such as non-melanoma skin cancers.²⁸ Here we have demonstrated for the first time that Δ Np63 α negatively regulates PTEN expression, and subsequently enhances Akt activation, creating a positive feedback loop that regulates keratinocyte proliferation, and perturbation of this axis might lead to epithelial cancers.

It has been shown earlier that head and neck SCCs are reliant on high levels of Δ Np63 α for survival through repression of p73-mediated apoptosis.¹⁹ Our studies, however, show that in at least epidermal cells TAp73 has little role in the transcriptional regulation of PTEN in the absence of genotoxic stress (Supplementary Figure S3). Previous studies demonstrated that TAp73 can induce PTEN transcription, suggesting that regulation of PTEN by the different p53 family members may be tissue specific.²⁹ Furthermore, in our studies we failed to detect p63 binding to the same region as p73 (region D), suggesting that p63 and p73 control PTEN expression through different REs.²⁹ Both TAp73 and TAp63 do not seem to induce PTEN expression, suggesting that Δ Np63 α -mediated PTEN repression is independent of TAp63 and TAp73 (Supplementary Figure S3).

Although p53 is capable of inducing the transcription of PTEN, Δ Np63 α functions as a master regulator independently of p53 to repress PTEN, as increases in PTEN expression were observed in A431 and HaCaT cells, which harbor different p53 mutations (R273H and p53Y179/W282, respectively), and in primary human epidermal keratinocytes (HEKs)

with wild-type p53. Our results wherein we observed that p53 bound to many of the same regions within the *pten* promoter where p63 binds (Supplementary Figure S4) could be explained by a previous report that has shown that mutant p53 can bind to p63.²⁰ Luciferase assays carried out in the absence of any p53 in cells overexpressing Δ Np63 α still show repression of PTEN-Luc activity (Figure 2c), suggesting that although it is possible for mutant p53 to bind Δ Np63 α , it does not alter the effects of p63 on PTEN regulation. It would be interesting to examine whether the differences in p53 mutational status affected other p63 target genes or whether the different p53 mutations dictate the tumorigenic and non-tumorigenic natures of A431 and HaCaT cells, respectively.

Mutations in the PI3K/Akt pathway are common in many types of cancers and contribute to the highly proliferative nature of tumors. Although overexpression of both Δ Np63 α and CA-Akt did not lead to a synergistic or additive decrease in PTEN expression, it did result in a larger decrease in PTEN expression compared with CA-Akt alone (Figure 4b). Δ Np63 α is more capable of repressing PTEN expression than CA-Akt alone, possibly due to the ability of active Akt to initiate the ARF/Egr-1 axis that induces PTEN to serve as a natural negative feedback mechanism.³⁰ The potent ability of Δ Np63 α to reduce PTEN expression suggests that Δ Np63 α amplification may be enough to tip the balance toward hyperplasia and cancer development. Δ Np63 α may also be capable of influencing proliferation by controlling nuclear levels of PTEN. Nuclear PTEN can elicit cell-cycle arrest independently of Akt, in part by binding to the anaphase-promoting complex to enhance cell-cycle repression.^{13,31} Our results in Figure 5 show that modulation of Δ Np63 α predominantly affects nuclear PTEN, supporting the hypothesis that Δ Np63 α can control proliferation in both an

Akt-dependent and -independent manner by repressing *pten* gene expression and by regulating PTEN localization, respectively. It remains to be seen, however, how Δ Np63 α is capable of regulating the nuclear import and/or export of PTEN. Given that little nuclear PTEN is observed in basal layer keratinocytes of normal skin, it is unlikely that the lack of nuclear PTEN seen in BCC and SCC could be used as a prognostic marker (Figure 6). Instead, the dramatic lack of nuclear PTEN in both BCC and SCC, presumably due in part to amplification of Δ Np63 α , might prevent cells from undergoing a proper cell-cycle arrest program.

As minute reductions in PTEN are sufficient to drive tumorigenesis, it is possible that changes in proteins that regulate PTEN, such as Δ Np63 α , could also contribute to tumor formation.¹⁷ We observed differences in the magnitude of proliferation after silencing PTEN or Δ Np63 α between cell lines (Figure 4b). A431 cells were less susceptible to changes in growth rates, possibly due to the atypical nature of this SCC line. However, both A431 and HaCaT cells showed similar trends in that silencing both Δ Np63 α and PTEN resulted in nearly normal growth rates, suggesting that both cell lines utilize the Δ Np63 α /PTEN axis to coordinate cell proliferation. The importance of the balance between Δ Np63 α and PTEN is further highlighted in SCC and BCC, the two most common cancers, where the ratio of p63 to PTEN is significantly perturbed (Figure 6). BCCs often display wild-type p53 and rarely invade into the surrounding tissue, whereas SCCs usually have mutated p53 and a higher propensity for invasion.^{32,33} The amplification of Δ Np63 α and the subsequent imbalance between Δ Np63 α and PTEN in both BCC and SCC suggests that regulation of PTEN in keratinocytes relies primarily on Δ Np63 α to regulate proliferation, and disruption of the Δ Np63 α /PTEN axis is an early insult contributing to both forms of skin cancer. Together, this study demonstrates that Δ Np63 α maintains normal cell proliferation by negatively regulating basal PTEN levels and that disruption of this balance could potentiate cell growth via activation of the Akt pathway, as well as through repression of nuclear PTEN.

Materials and Methods

Cell lines and transfections. H1299, A431, HaCaT, PC3 and E1A-transformed primary mouse embryonic fibroblasts (gifted by Dr. Elsa Flores, MD Anderson) were maintained as described earlier.³⁴ Primary human epidermal keratinocytes were maintained in Epilife media supplemented with human keratinocyte growth supplement and 100 U/ml penicillin and streptomycin and 0.25 μ g/ml amphotericin B (Cascade Biologics, Portland, OR, USA). The Δ Np63 α and CA-Akt plasmids used were described previously.^{34,35} H1299 cells were transfected using Lipofectamine 2000, and A431, HaCaT and HEK cells were transfected with two rounds of siRNA using Oligofectamine as per the manufacturer's instructions (Invitrogen, Carlsband, CA, USA). Cells were harvested for immunoblot and qRT-PCR studies 24 h after the second transfection. Three different siRNAs (sip63-1, sip63-2 and sip63-3) were used to silence p63 to rule out off-target effects. Sequences used for silencing total p63, TAp63 and Δ Np63 were described previously.^{34,36,37} Sequences for silencing TAp73 were described previously.³⁸ Sequences used for silencing PTEN are sense r(5'-GGCGUUAACAGGAACAAUA-3')dTdT and antisense r(5'-UAUUGUUCUGU AUACGCC-3')dTdT.

RNA isolation and TaqMan real-time PCR studies. Quantitative real-time PCR analysis was carried out as previously described using Assay on Demand (AOD) specific for genes of interest and normalized to endogenous GAPDH for human genes or to β -actin for murine genes of interest (PE Applied Biosystems,

Foster City, CA, USA).³⁴ Human AODs used were GAPDH (4325792), p63 (Hs_00978340_m1), Δ Np63 (Hs_00978339_m1), TAp63 (Hs_00186613_m1), TAp73 (Hs_01056228_m1), Δ Np73 (Hs_1065727) IGFBP-3 (Hs_00426287_m1) and PTEN (Hs_00829813_s1). Murine AODs were p63 (Mm00495788_m1), PTEN (Mm00477210_m1) and β -actin (Mm00607939_s1).

PTEN transactivation assays. The full-length PTEN-luc reporter used was described previously.³⁹ Cells were transfected with 100 ng of reporters containing the full-length PTEN promoter or 500 ng of the designated minimal PTEN promoter reporter along with the Δ Np63 α or empty vector and *Renilla* luciferase expression vector to control for transfection efficiency. Dual-luciferase assays were carried out using Dual-Luciferase Reporter 1000 Assay System as per manufacturer's protocol (Promega, Madison, WI, USA). The minimal PTEN luciferase reporters were created by annealing primers to the regions of interest followed by ligation into the pGL3 Basic promoter-less luciferase reporter (Promega) at the KPM and HINDIII sites. The primers used were: A forward 5'-CAACGAGCCGAGTTACCGGG AAGCGAGAGGTGGGCGCTGCAAGGGAA-3', A reverse 5'-AGCTTCCCTTG CAGCGCCCCACCTCTCGCTTCCCGGTAACCTCGGCTCGTTTGGTAC-3'; B forward 5'-CCAGCGTGGTCACTGCTGCTTTTACCTGTGCACAGGTAACCTCAG ACTCGAGTCAA-3', B reverse 5'-AGCTTTGACTCGAGTCTGAGGTTACCTGTGC ACAGGTGAAAAGGACAGGTGACCAAGCTGGGTAC-3'; C forward 5'-CGAGC AAGCCCCAGGCAGCTACACTGGGCATGCTCA-3', C reverse 5'-AGCTTGAGCA TGCCCAAGTGTAGCTGCCTGGGGCTTGCTCGGTAC-3'; and E forward 5'-CAAC CGTGCAGCCTCTTCTCGGCTTCTCTGAAAGGGAAGGTGGAAGCCGTGGG CA-3', E reverse 5'-AGCTTGCCACGGCTCCACCTTCCCTTCAGGAGAAGC CGAGGAAGAGGCTGCACGGTTGGTAC-3'.

Chromatin immunoprecipitation assays (ChIP). Chromatin immunoprecipitation was carried out using the ChIP-IT Express Kit as per the manufacturer's protocol (Active Motif, Carlsbad, CA, USA) and as described previously.³⁴ Sequences for primers are PTEN A forward 5'-CGGGGTACCCGTCC CACTCAGGGATCCTCT-3' and reverse 5'-CCCAAGCTTCTGCTATTGTGTCG CCAGCGTCT-3'; PTEN B forward 5'-CCCTGCATTTCTCTACA-3' and reverse 5'-GAGGCGAGGATAACGAGCTA-3'; PTEN C forward 5'-GCACCATCTCAGC TTTCAT-3' and reverse 5'-CAAGCCGAGGCTCTACT-3'; PTEN D forward 5'-GA CTGCGCTCAGTTCTCTCC-3' and reverse 5'-AGAGGCTGCACGGTTAGAAA-3'; and PTEN E forward 5'-CGGGGTACCAGCGCGCGCTTCTCGCCTCT-3' and reverse 5'-CCCAAGCTTAGACCCCTCCCTGGAGCTGCAG-3'; primers for the p21 promoter were as described previously.³⁴ PCR conditions used are as follows: a total of 40 cycles were performed, each consisting of 30 s at 94°C, 30 s at 55°C and 30 s at 68°C.

Western blot analysis. Whole-cell extracts were prepared by lysing the cells in phosphatase inhibitors containing buffer (50 mM Tris-HCl pH 8, 120 mM NaCl, 5 mM NaPPi, 10 mM NaF, 30 mM paratitrophenylphosphate, 1 mM benzamide, 0.1% NP-40, 1% Triton X-100 and 0.2 mM PMSF, 100 nM sodium orthovanadate) supplemented with protease inhibitors. Subcellular fractionations were carried out using the NE-PER kit as per the manufacturer's instructions (Pierce, Rockford, IL, USA). Immunoblot analysis was carried out as described previously.³⁴

Antibodies. p63 antibodies used are as follows: pan-p63 4A4, p63alpha-isoform-specific H-129 (Santa Cruz Biotechnology, Inc., Santa Cruz, CA, USA), and Δ Np63-specific RR-14 and TAp63-specific antibodies (courtesy of Dr. Satrajit Sinha, SUNY Buffalo).⁴⁰ The p53 antibody (Ab-6) was purchased from Calbiochem (Gibbstown, NJ, USA). PTEN (#9552), phospho-Akt (Ser 473, #9271), total Akt (#9272) and total PARP (#9542) antibodies were purchased from Cell Signaling (Danvers, MA, USA), whereas monoclonal anti- β -actin was purchased from Sigma Aldrich (St. Louis, MO, USA). Antibodies specific to total p73 (IMG-259A) and Δ Np73 (IMG-313A) were purchased from Imgenex (San Diego, CA, USA). The antibody for α -tubulin was graciously donated by Dr. Paula Bubulya (Wright State University). Appropriate horseradish peroxidase-conjugated secondary antibodies (Promega) were used for chemiluminescence detection with Supersignal Westpico Chemiluminescent Substrate kit (Pierce).

Cell growth assays. Cells harvested after a second round of siRNA transfection were re-seeded at 2×10^4 cells/well in a six-well plate for colony-formation assays. At 72 h, cells were stained with 1% crystal violet. A total of 20 random images were taken per condition, and colony number from each frame was measured using ImagePro 6.2 Software (Media Cybernetics, Bethesda, MD, USA).

For cell growth assays, cells were re-seeded at 5000 cells/well in 96-well-plate MTS assays or at 10 000 cells/well in 12-well plates for trypan blue exclusion assays after 24 h from the second round of siRNA transfection. The MTS assay was carried out as per the manufacturer's instructions at the days indicated (Promega). Cell proliferation as measured by trypan blue exclusion was carried out in triplicate per condition per day post siRNA transfection.

Immunofluorescence. Human skin microarray (SK805, Biomax, Rockville, MD, USA) and paraffin-embedded embryonic mouse sections (a kind gift from Dr. Satrajit Sinha, SUNY Buffalo) were stained as previously described with minor modifications.³⁴ Modifications included an antigen retrieval of 12 min in Tris/EDTA buffer (10 mM Tris, 1 mM EDTA, 0.05% Tween 20, pH 9.0) at 90°C in a pressure cooker and image was captured using a Leica CTR 6000 microscope (Leica Microsystems, Wetzlar, Germany) with ImagePro 6.2 Software (Media Cybernetics). AlexaFluor 488 and 568 secondary antibodies were used to visualize p63 and PTEN, respectively (Invitrogen). MFI for each tissue sample was calculated using ImagePro 6.2 software after normalization for background intensity. Multiple measurements (at least 5), all of the same size, were taken of the epidermal tissue for each tissue sample. All experimental procedures involving mice were approved by the Institutional Animal Care and Use Committee of the State University of New York at Buffalo.³⁴ Human tissue samples were obtained and handled in full approval and accordance with Wright State University IRB protocols.

Statistical analyses. Statistical significance of differences was determined by Student's *t*-test for all experiments except MFIs, where the Pearson's product moment analysis was used to determine correlations between p63 and PTEN and significant difference was determined by one-way ANOVA with a Bonferroni's *ad-hoc* test. A probability value of <0.05 was considered statistically significant.

Conflict of Interest

The authors declare no conflict of interest.

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Supplementary Information accompanies the paper on Cell Death and Differentiation website (<http://www.nature.com/cdd>)