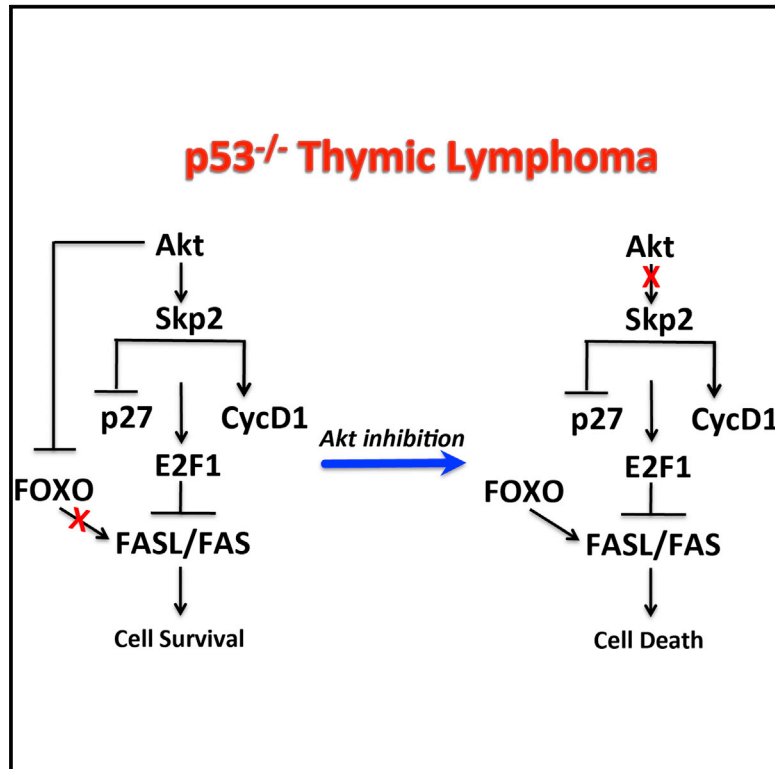**Article**

Systemic Akt1 Deletion after Tumor Onset in $p53^{-/-}$ Mice Increases Lifespan and Regresses Thymic Lymphoma Emulating p53 Restoration

Graphical Abstract

Authors

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In Brief

Yu et al. have deleted Akt1 after tumor onset in $p53^{-/-}$ mice. Systemic Akt1 deletion regresses thymic lymphoma in $p53^{-/-}$ mice emulating p53 restoration. Furthermore, pharmacological inhibition of Akt selectively kills thymic lymphoma cells and not primary thymocytes.

Highlights

- Systemic Akt1 deletion regresses $p53^{-/-}$ thymic lymphoma, emulating p53 restoration

- Akt inhibition regresses tumors that are not driven by Akt activation

- Skp2 inhibits apoptosis by inhibiting FasL and Fas expression

- Overexpression of Skp2 could exert resistance to Akt inhibition

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Systemic Akt1 Deletion after Tumor Onset in *p53*^{−/−} Mice Increases Lifespan and Regresses Thymic Lymphoma Emulating p53 Restoration

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SUMMARY

Akt is frequently activated in human cancers. However, it is unknown whether systemic inhibition of a single Akt isoform could regress cancer progression in cancers that are not driven by Akt activation. We systemically deleted Akt1 after tumor onset in *p53*^{−/−} mice, which develop tumors independently of Akt activation. Systemic Akt1 deletion regresses thymic lymphoma in *p53*^{−/−} mice emulating p53 restoration. Furthermore, pharmacological inhibition of Akt selectively kills thymic lymphoma cells and not primary thymocytes. Mechanistically, Akt1 inhibition in *p53*^{−/−} thymic lymphoma inhibits Skp2 expression and induces FasL, which is the primary cause of cell death. Skp2 exerts resistance to cell death by antagonizing the induction of FasL and reducing FAS expression, which is linked to cyclin D1 expression. The results established a paradigm whereby systemic Akt1 inhibition is sufficient to regress tumors that are not driven by Akt activation and a mechanism of cell survival by Skp2.

INTRODUCTION

The serine/threonine kinase Akt is perhaps the most frequently activated oncoprotein in human cancers (reviewed in Bhaskar and Hay, 2007). The pathways leading to its activation as well as Akt itself are being targeted for cancer therapy. However, there are three Akt isoforms (Akt1–3) encoded by three separate genes. These isoforms are about 85% identical to each other, and it is impossible to distinguish among them with respect to substrate specificity in in vitro kinase assay (Walker et al., 1998). Nevertheless, it is possible that the different isoforms are functionally different in vivo. However, it is unknown which isoform should be preferentially targeted and for which type of cancer, without eliciting adverse physiological consequences. It is also not known whether Akt could be targeted for therapy of cancers that are not driven directly or indirectly by Akt activation.

Previously, we and others showed that germline deletion of Akt1 exerts resistance to Pten-deletion-induced tumorigenesis (Chen et al., 2006), skin carcinogenesis (Skeen et al., 2006), mammary and salivary gland tumorigenesis (Ju et al., 2007; Maroulakou et al., 2007; Skeen et al., 2006), and lung carcinogenesis (Hollander et al., 2011). The germline deletion of Akt1 in the mouse does not have adverse physiological consequences, and *Akt1*^{−/−} mice live significantly longer than control WT mice (Chen et al., 2006). By contrast, germline deletion of Akt2 induces insulin resistance and does not inhibit tumor development in mouse models of breast cancer and in Pten^{+/−} mice (Maroulakou et al., 2007; Xu et al., 2012). However, studies using germline deletions of Akt isoforms cannot determine whether Akt is required for tumor progression and do not emulate drug therapy. To address whether systemic Akt1 deletion after tumor onset could inhibit tumor progression even in tumors, which are not driven by Akt activation, we examined the ability of systemic Akt1 deletion to inhibit tumor progression in *p53*^{−/−} mice. We found that the systemic deletion of Akt1 was sufficient to regress or halt the progression of thymic lymphoma in *p53*^{−/−} mice similar to what observed after p53 restoration in these mice (Ventura et al., 2007). The deletion of Akt1 in *p53*^{−/−} thymic lymphoma is sufficient to inhibit cell-cycle progression and induces apoptosis, which could be recapitulated by a pharmacological inhibitor of Akt. The cytotoxic effect of Akt inhibition on *p53*^{−/−} thymic lymphoma is selective as it is not cytotoxic to normal thymocytes. However, hyperactivation of Akt in p53^{−/−} thymic lymphoma exerts resistance to pharmacological inhibitor of Akt. The induction of FasL expression by the inhibition of Akt and activation of FOXO appears to be the predominant cause of cell death. Interestingly ectopic expression of Skp2 in the *p53*^{−/−} thymic lymphoma cells also exerts resistance to Akt inhibition. Skp2 exerts resistance to cell death at least in part by the inhibition of FasL and Fas expression. It was recently reported that D cyclins promote survival of hematopoietic cells through inhibition of Fas and FasL expression in a E2F1-dependent manner (Choi et al., 2014). Consistently, we found that the ectopic expression of Skp2 also elevates Cyclin D1 expression and reduces p27 expression, which are known to activate E2F1, and thus could explain the mechanism by which Skp2 inhibits Fas and FasL expression. Taken together, these results established a paradigm whereby Akt1 inhibition is therapeutic
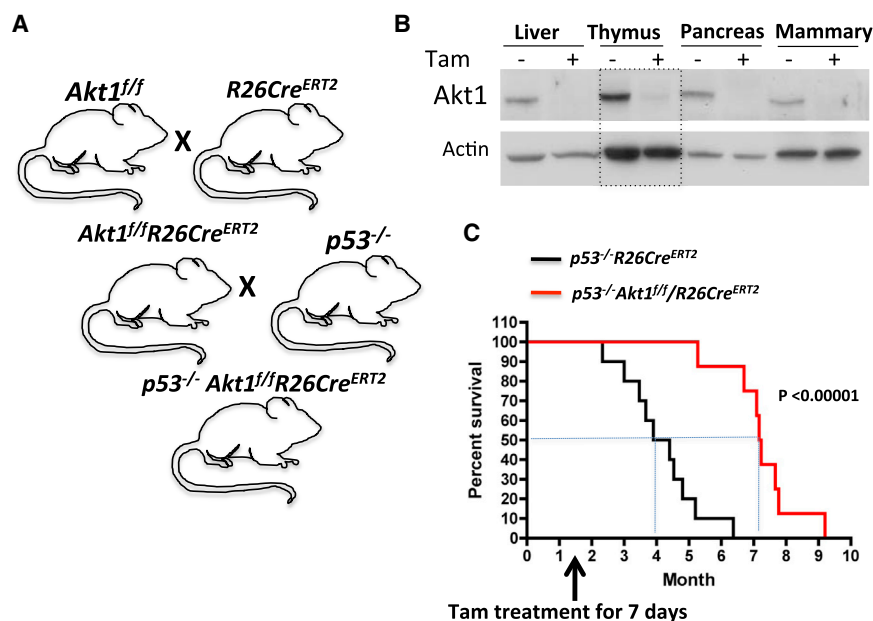
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A

$Akt1^{f/f}$ X $R26Cre^{ERT2}$

$Akt1^{f/f}R26Cre^{ERT2}$ X $p53^{-/-}$

$p53^{-/-} Akt1^{f/f}R26Cre^{ERT2}$

B

| | Liver | | Thymus | | Pancreas | | Mammary | |
Tam: − + − + − + − +
Akt1
Actin

C

— $p53^{-/-}R26Cre^{ERT2}$
— $p53^{-/-}Akt1^{f/f}/R26Cre^{ERT2}$

P <0.00001

Percent survival vs Month

Tam treatment for 7 days

Figure 1. Systemic Deletion of Akt1 in $p53^{-/-}$ Mice after Tumor Onset Increased Lifespan
(A) Schematic illustration showing crossing of mice to generate $p53^{-/-}Akt1^{f/f}/R26Cre^{ERT2}$ mice.
(B) Immunoblot showing Akt1 deletion in several mouse tissues after injection of tamoxifen (Tam) to 6-week-old $p53^{-/-}Akt1^{f/f}/R26Cre^{ERT2}$ mice.
(C) Kaplan-Meier plot showing lifespan of $p53^{-/-}$ or $p53^{-/-}Akt1^{f/f}/R26Cre^{ERT2}$ mice after tamoxifen injection at 6 weeks of age (n = 10).

even for cancers that are not driven by Akt activation. These results suggest that Akt inhibitors could be used effectively to treat cancers that are driven by the loss of p53 such as in Li-Fraumeni syndrome (LFS) patients. Although only about 20% of LFS patients have a compete loss of p53, most of the p53 mutations in LFS are occurring in the DNA binding domain and act in a dominant manner to inhibit WT p53 (Malkin, 2011). The results also suggest that high Skp2 expression could predict resistance to Akt inhibition and that the inhibition of both Akt and Skp2 might be considered for a long-term therapeutic effect.

RESULTS

Systemic Deletion of Akt1 in $p53^{-/-}$ Mice Increases Lifespan and Halts the Progression of Thymic Lymphoma in These Mice

To determine the ability of systemic Akt1 ablation to regress or attenuate tumor progression, we have generated $Akt1^{f/f}$ mice (Figures S1A and S1B) and crossed these mice with $R26Cre^{ERT2}$ mice, in which CreERT2 was inserted in the ubiquitously expressed ROSA26 locus (Ventura et al., 2007), to generate $Akt1^{f/f}/R26Cre^{ERT2}$ mice (Figures 1A and S1C). CreERT2 is an inducible Cre recombinase fused in frame to a mutated ligand-binding domain of estrogen receptor, which responds only to hydroxytamoxifen (4-OHT) and not to estrogen. To determine whether systemic Akt1 inhibition could regress or inhibit progression of tumors that are not driven by lesions that activate Akt, we crossed $Akt1^{f/f}/R26Cre^{ERT2}$ mice with $p53^{-/-}$ mice to generate $p53^{-/-}Akt1^{f/f}/R26Cre^{ERT2}$ mice (Figure 1A). Injection of tamoxifen to the mice 6 weeks after birth successfully deletes Akt1 in all tissues tested (Figures 1B and S1). The $p53^{-/-}$ mice develop thymic lymphoma with an early onset and eventually develop also sarcoma (Ventura et al., 2007). The mice die of these tumors within 6 months after birth, and restoration of p53 regresses the lymphoma with increased apoptosis and inhibits the growth

of sarcoma, indicating that these tumors are highly dependent on p53 deletion (Ventura et al., 2007). We have generated cohorts of $p53^{-/-}Akt1^{f/f}/R26Cre^{ERT2}$ and $p53^{-/-}R26Cre^{ERT2}$ mice. The mice were injected with tamoxifen for 7 consecutive days, 6 weeks after birth when they start developing thymic lymphoma. As shown in Figure 1C, the systemic deletion of Akt1 substantially increased the median lifespan of $p53^{-/-}$ mice from 16 weeks to 29 weeks. We did not see a significant change in the lifespan of control $p53^{-/-}R26Cre^{ERT2}$ mice (data not shown).

To further substantiate our findings and demonstrate that systemic Akt1 ablation is sufficient to regress or halt thymic lymphoma progression in $p53^{-/-}$ mice, we monitored in vivo tumor progression by MRI prior to and after treatment with tamoxifen. Cohorts of $p53^{-/-}Akt1^{f/f}/R26Cre^{ERT2}$ and $p53^{-/-}R26Cre^{ERT2}$ mice were monitored to detect in vivo lymphoma as previously described (Ventura et al., 2007). When tumors detected by MRI reached the volume of approximately 200 mm^3, mice were treated with tamoxifen, and tumor growth was followed as previously described (Ventura et al., 2007). As shown in Figures 2A and 2B, tumor size was significantly reduced 12–16 days after the administration of tamoxifen into $p53^{-/-}Akt1^{f/f}/R26Cre^{ERT2}$ mice, while at the same time in untreated mice tumors grew approximately by about 5-fold (Figure 2A). Injection of tamoxifen into control $p53^{-/-}R26Cre^{ERT2}$ mice did not reduce tumor growth (Figure 2B). In general, the deletion of Akt1 by tamoxifen treatment either halts or regresses tumor growth while at the same time in control mice tumor grew by 2- to 6-fold (Figure 2C). Moreover, analysis of sections from the tumor mass that was left after Akt1 deletion revealed a massive apoptosis as indicated by cleaved caspase-3 staining and a relatively few Ki67 positive cells (Figure 2D). These results indicate that systemic deletion of Akt1 after tumor onset in $p53^{-/-}$ mice, which emulate drug therapy, is sufficient to regress $p53^{-/-}$ thymic lymphoma. Moreover, the effect of Akt1 deletion on thymic lymphoma in $p53^{-/-}$ mice is similar to what observed after restoration of p53 in these mice (Ventura et al., 2007).

The Deletion of Akt1 in p53$^{-/-}$ Thymic Lymphoma Cells Induces Cell Death and Inhibits Cell Proliferation

To further establish the role of Akt1 in p53$^{-/-}$ thymic lymphoma, cell lines were isolated from thymic lymphoma tumors in
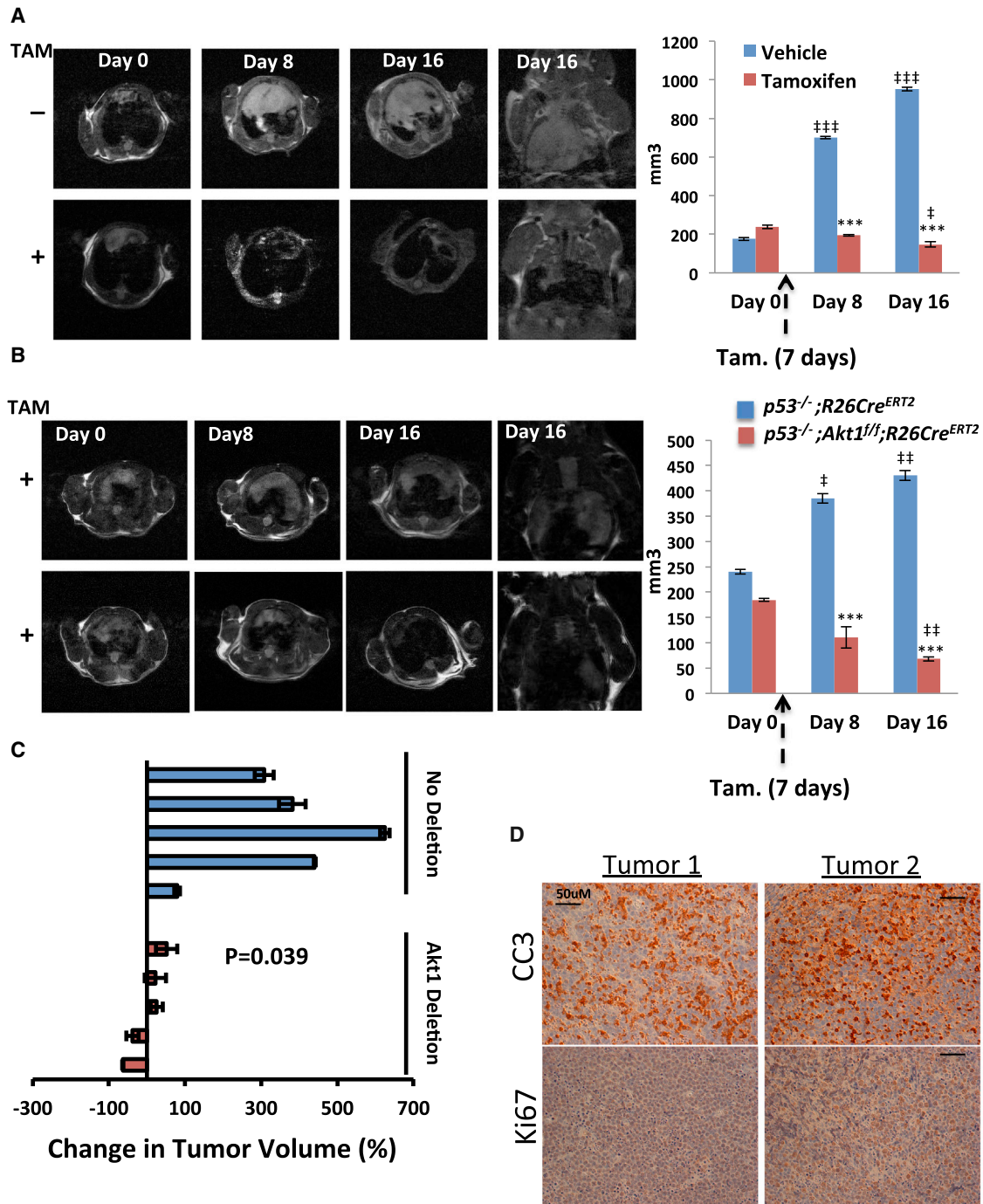
Figure 2. Systemic Akt1 Deletion in *p53−/−* Mice Halts or Regresses Tumor Growth In Vivo

(A and B) MRI images (left) and calculated tumor volume (right) in response to tamoxifen of *p53−/−Akt1f/f/R26CreERT2* (A and B) or *p53−/−R26CreERT2* mice (B). ***p < 0.001 Tam versus vehicle; ‡‡‡p < 0.0001, ‡‡ p < 0.005, ‡ p < 0.01 versus day 0.

(C) Summary of individual tumor responses to systemic Akt1 deletion 16 days after tamoxifen treatment or vehicle alone of *p53−/−Akt1f/f/R26CreERT2* mice.

(D) Representative tumor sections from two tumors after regression as determined by MRI showing cleaved caspase-3 (CC3) and Ki67 staining. Data are presented as an average of three measurements ±SEM.

p53−/−R26CreERT2 and *p53−/−Akt1f/f/R26CreERT2* mice and were exposed to 4-OHT in vitro (Figure 3A). The deletion of Akt1 in these lymphoma cell lines profoundly inhibited the growth

of the cells (Figure 3B). This effect is due to both induction of apoptosis and inhibition of proliferation as measured by bromodeoxyuridine (BrdU) incorporation (Figure 3C). Thus, Akt1
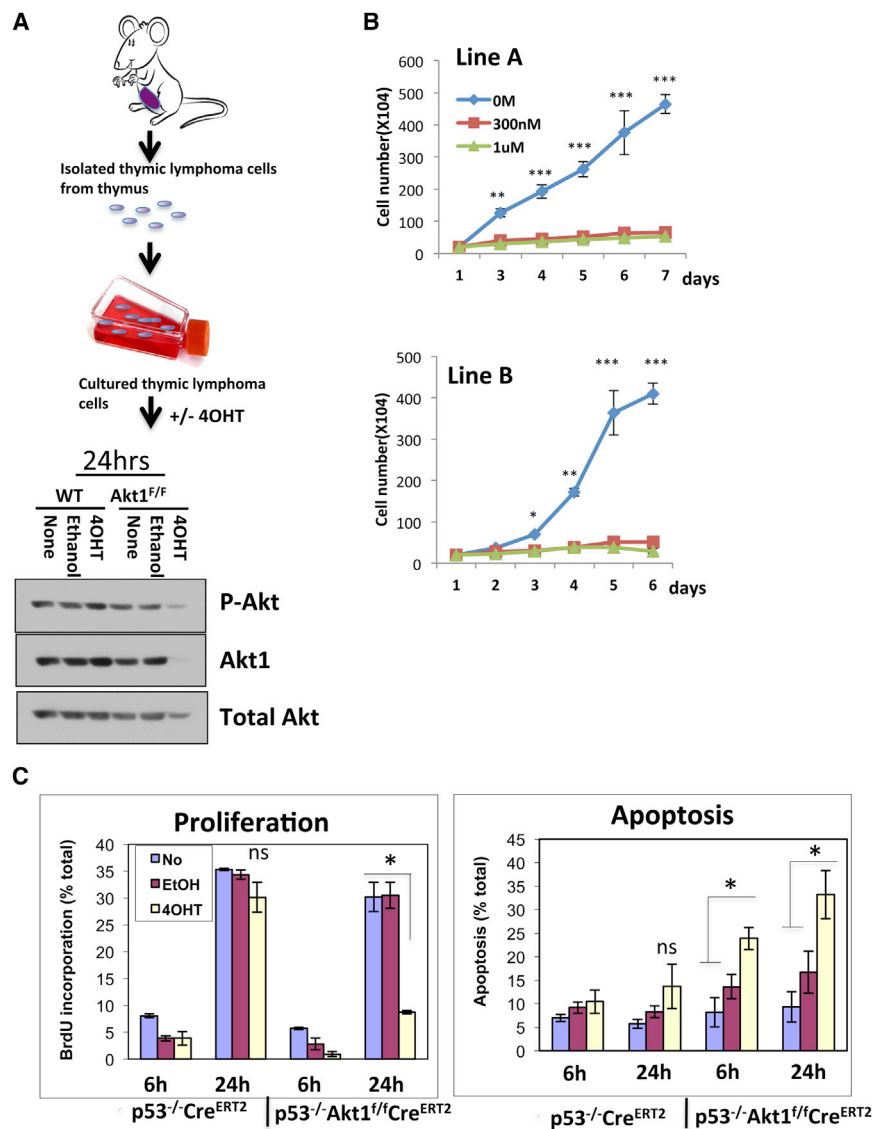
Figure 3. Ex Vivo Deletion of Akt1 in Thymic Lymphoma Cells Derived from $p53^{-/-}$ $Akt1^{f/f}/R26Cre^{ERT2}$ Mice Inhibits Proliferation and Induces Cell Death

(A) Strategy of isolating thymic lymphoma cell lines (top) and immunoblot showing Akt1 deletion after exposure to 4-OHT (bottom).

(B) Proliferation of two independent isolated cell lines in response to 4-OHT. Cells were exposed to 4-OHT for 3 days, and cell number was determined in the indicated time points. Data are presented as an average of three independent experiments in triplicates ±SEM. ***p < 0.001, **p < 0.01, *p < 0.05 versus 4-OHT 300 nM and 1 μM.

(C) BrdU (bromodeoxyuridine) incorporation as measured by anti-BrdU staining and cell death as measured by PI (propidium iodide) staining using flow cytometry. Cells were exposed to 4-OHT for 3 days and were subjected to BrdU incorporation and PI staining at the indicated time points. Data are presented as an average of three independent experiments in triplicates ±SEM. *p < 0.05; ns, not significant.

deletion is sufficient to induce both cell death and cell-cycle arrest in p53$^{-/-}$ thymic lymphoma. In general, Akt1 deletion by itself has only cytostatic effect and can only render cells more susceptible to cell death by the presence of other apoptotic stimuli (Chen et al., 2001). Notably, the p53$^{-/-}$ thymocytes do not display activation of Akt when compared to control thymocytes, and Akt activity did not significantly change after re-expressing p53 in the p53$^{-/-}$ thymic lymphoma cells (Figure S2). Thus, clearly Akt is not hyperactivated in thymic lymphoma driven by p53 deletion. Ectopic expression of a myristoylated Akt1 (mAkt), a constitutively active form of Akt, exerts resistance to the deletion of Akt1 by 4-OHT (Figures S2C and S2D). Finally, we injected either p53$^{-/-}$/R26Cre^{ERT2} or p53$^{-/-}$Akt1$^{f/f}$/R26Cre^{ERT2} lymphoma cells into nude mice and followed tumor growth (Figure 4A). When tumor size reached 60 mm^3 in volume, the mice were treated with tamoxifen. While tumor growth of p53$^{-/-}$/R26Cre^{ERT2} lymphoma cells was not affected by tamox-

ifen treatment, tumor growth of p53$^{-/-}$ Akt1$^{f/f}$/R26Cre^{ERT2} lymphoma cells in the treated mice was halted when compared to tumor growth of untreated mice (Figures 4B and 4C). When, tumor section samples were analyzed for cell proliferation or cell death as measured by BrdU incorporation and cleaved caspase-3, respectively, a profound reduction in proliferation and increased apoptosis were observed (Figure 4D). Thus, the in vivo results recapitulate the results obtained ex vivo.

We then wanted to find out whether pharmacological inhibition of Akt1 could recapitulate Akt1 deletion. Since there are no specific inhibitors of Akt1, we employed MK2206, which is an allosteric inhibitor of Akt that is commonly used in clinical trials. MK2206 inhibits Akt1, Akt2, and Akt3 with IC$_{50}$ of 8, 12, and 65 nM, respectively (Hirai et al., 2010). We first exposed the thymic lymphoma cells to MK2206 and compared its effect to the effect induced by Akt1 deletion. We found a similar effect of MK2206 on the proliferation of the cells (Figure 5A) as well on BrdU incorporation (Figure 5B) and apoptosis (Figure 5C) to that observed after the induction of Akt1 deletion. Likewise in vivo tumor growth of these cells was halted after treatment with MK2206 (Figure 5D) similar to what was observed after systemic deletion of Akt1. Thus, the pan-Akt inhibitor could recapitulate the conditional deletion of Akt1 to inhibit the progression of cancer that is not driven by oncogenic signaling that activates Akt. Importantly, the effect of MK2206 on p53$^{-/-}$ thymic lymphoma is selective because we found that the cytotoxic effect on primary normal thymocytes is minimal in comparison to the lymphoma cells (Figure 5E).
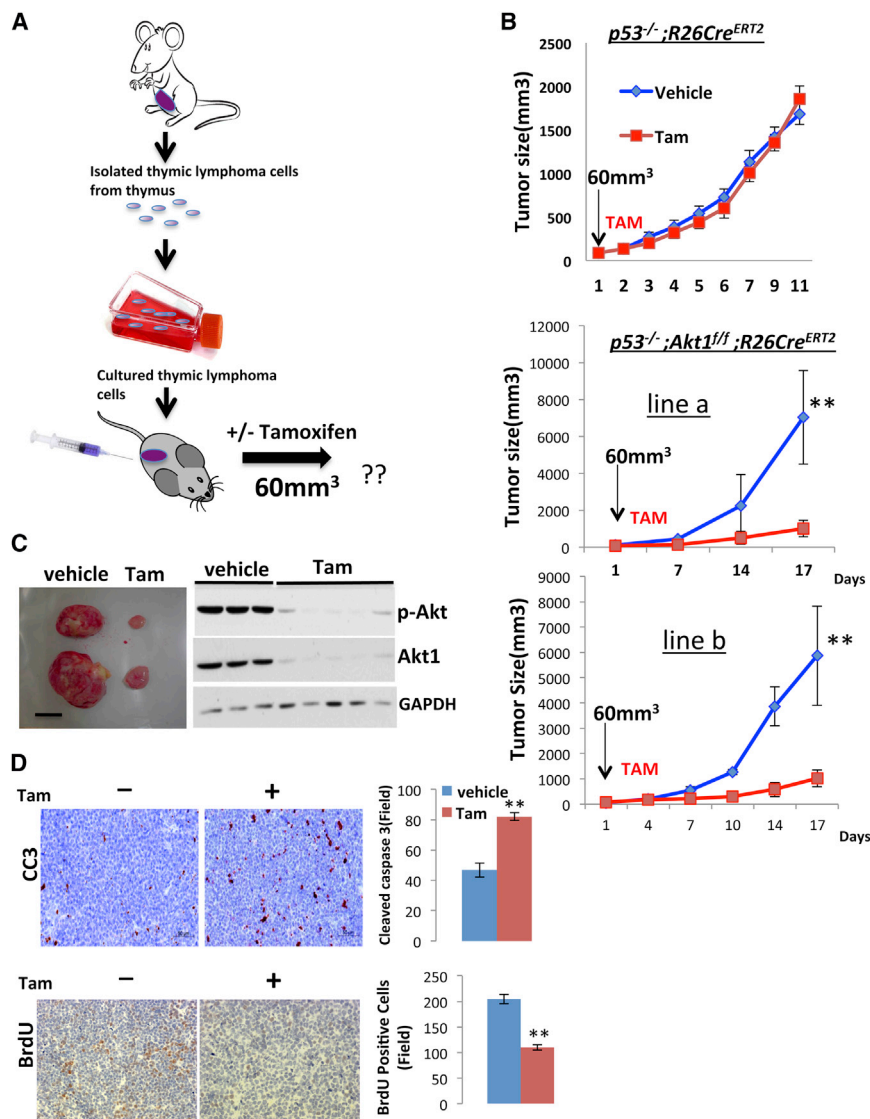
Figure 4. The Deletion of Akt1 Inhibits the Growth of Xenographic p53$^{-/-}$ Thymic Lymphoma Tumors by Inhibiting Proliferation and Inducing Apoptosis

(A) Flowchart depicting the strategy to determine tumor response in xenograft assays. Cells isolated from the thymic lymphoma were cultured and inoculated into nude mice. When tumor size reached 60 mm^3, mice were treated with tamoxifen, and tumor growth was determined.

(B) Tumor growth curves of *p53$^{-/-}$R26Cre^{ERT2}* or *p53$^{-/-}$Akt1$^{f/f}$/R26Cre^{ERT2}* cells in response to tamoxifen. Data are presented as an average of ten tumors in five mice per each cell line and per each treatment ±SEM (n = 9). **p < 0.001 versus Tam.

(C) Representative tumors from vehicle- or tamoxifen-treated mice (left) and immunoblot showing the extent by which Akt1 deletion affected total serine 473 phosphorylation of Akt in various tumors (right).

(D) Representative tumor sections from vehicle- or tamoxifen-treated mice showing cleaved caspase-3(CC3) and BrdU incorporation (left panels). Quantification of CC3 or BrdU positive cells in randomly selected six fields from three different tumors (right panels). Data are presented as an average ±SEM. **p < 0.001.

exerted by p53 deficiency could be extended to different types of cancer.

The Mechanisms by which the Deletion of Akt1 Inhibits Proliferation and Induces Cell Death in p53$^{-/-}$ Thymic Lymphoma

We previously showed that mTORC1 is the most critical downstream effector of Akt, required for cell proliferation and tumorigenesis (Skeen et al., 2006). Our results showed that, upon deletion of Akt1 by 4-OHT, there was a decrease in FOXO phosphorylation (Figure 6A) and a marked reduction in mTORC1 activity as measured by the phosphorylation of S6 and 4EBP1 (Figure 6B). As we previously showed, mTORC1 activity and its downstream effector, the initiation factor of mRNA translation, eIF4E, are required for the Skp2 mRNA translation, and therefore for its expression at the protein level (Nogueira et al., 2012). Indeed, we found a significant decrease in Skp2 expression following the reduced mTORC1 activity (Figure 6B). Skp2 is a F-box protein that mediates the degradation of p27 and other cell-cycle inhibitory proteins (Nakayama et al., 2004; Pagano, 2004). Similarly, Akt inhibition by MK2206 inhibited FOXO phosphorylation and mTORC1 activity and elevated p27 expression (Figure 6C). Thus, the reduced mTORC1 activity in combination with the induced FOXO activity could explain the high level of p27 as well as the inhibition of proliferation following Akt inhibition in p53$^{-/-}$ thymic lymphoma. We also found a decrease in the anti-apoptotic protein Mcl-1 after Akt inhibition (Figure 6C). This is consistent with the high

Taken together, these results showed that Akt activity is also required for tumors that are not driven by signaling pathways that activate Akt, and that Akt ablation is therapeutic for thymic lymphoma induced by p53 deletion.

Interestingly, we previously showed that germline deletion of Akt1 sensitized primary thymocytes to killing by gamma-irradiation (Chen et al., 2001). Thus, it is possible that the deletion of p53, which increases DNA damage, similarly sensitizes thymocytes to killing by Akt inhibition. Consistently, we found that human cancer cells lacking p53 are profoundly more sensitive to cell death after inhibition of Akt then cancer cells proficient for p53. We employed three well-established pairs of colorectal cancer cell lines in which p53 is either WT or was somatically deleted (Sur et al., 2009; Figure S3). As shown in Figure 5F, the p53-deficient cells are highly sensitive to cell death after exposure to MK2206 when compared to the p53-proficient cells. These results suggest that the hypersensitivity to Akt inhibition

treated but only modestly in mAkt-treated cells, and the elevation of p27 was observed in control-treated cells but not in Skp2- or mAkt-treated cells (Figure 7B). Unlike in mAkt-treated cells, mTORC1 activity, as measured by S6K1 phosphorylation and 4EBP1 mobility shift, as well as FOXO phosphorylation are still inhibited to a similar extent in control and Skp2-ex-

dependency of Mcl-1 mRNA translation on mTORC1 activity and eIF4E (Wendel et al., 2007), as well as the increase of MCL-1 protein stability by Akt through the inhibition of GSK3 (Maurer et al., 2006). In addition, the mRNAs for the FOXO targets proapoptotic proteins Bim and FasL (Figures 6D and 7C), as well as FasL protein, which its expression, as determined by fluorescence-activated cell sorting (FACS) analysis, overlaps with cells undergoing cell death (Figures 6E and S4), were elevated after Akt inhibition. Thus, the reduced Mcl-1 expression together with the higher expression of Bim and FasL could explain the induction of cell death after Akt inhibition. Surprisingly, however, we found that ectopic expression of Skp2 in $p53^{-/-}$ thymic lymphoma exerted resistance to cell death induced by Akt inhibition almost to the same extent as by mAkt (Figure 7A). Analysis of protein extracts derived from these cells before and after treatment showed that Skp2 protein expression declined in control-

pressing cells. MCL-1 protein levels were reduced to a similar extent in control and Skp2-expressing cells (Figure 7B). Nonetheless, cleaved caspase-3 levels were profoundly reduced in Skp2 expression cells, consistent with the resistance of these cells to cell death induced by Akt inhibition. Thus, Skp2 could partially recapitulate the resistance to cell death exerted by activated Akt in p53$^{-/-}$ thymic lymphoma cells and overrides mTORC1 inhibition and FOXO activation. Therefore, the resistance to cell death induced by Skp2 overexpression seems to be independent of mTORC1 activity, FOXO phosphorylation, or Mcl-1 expression. These results were unexpected, but, although we did not observe a significant change in the induction of Bim mRNA in Skp2-expressing cells, we found that Fas ligand (FasL) expression was significantly reduced (Figure 7C). This could explain in part the resistance of p53$^{-/-}$ thymic lymphoma cells overexpressing Skp2 to Akt inhibition. To verify the role of
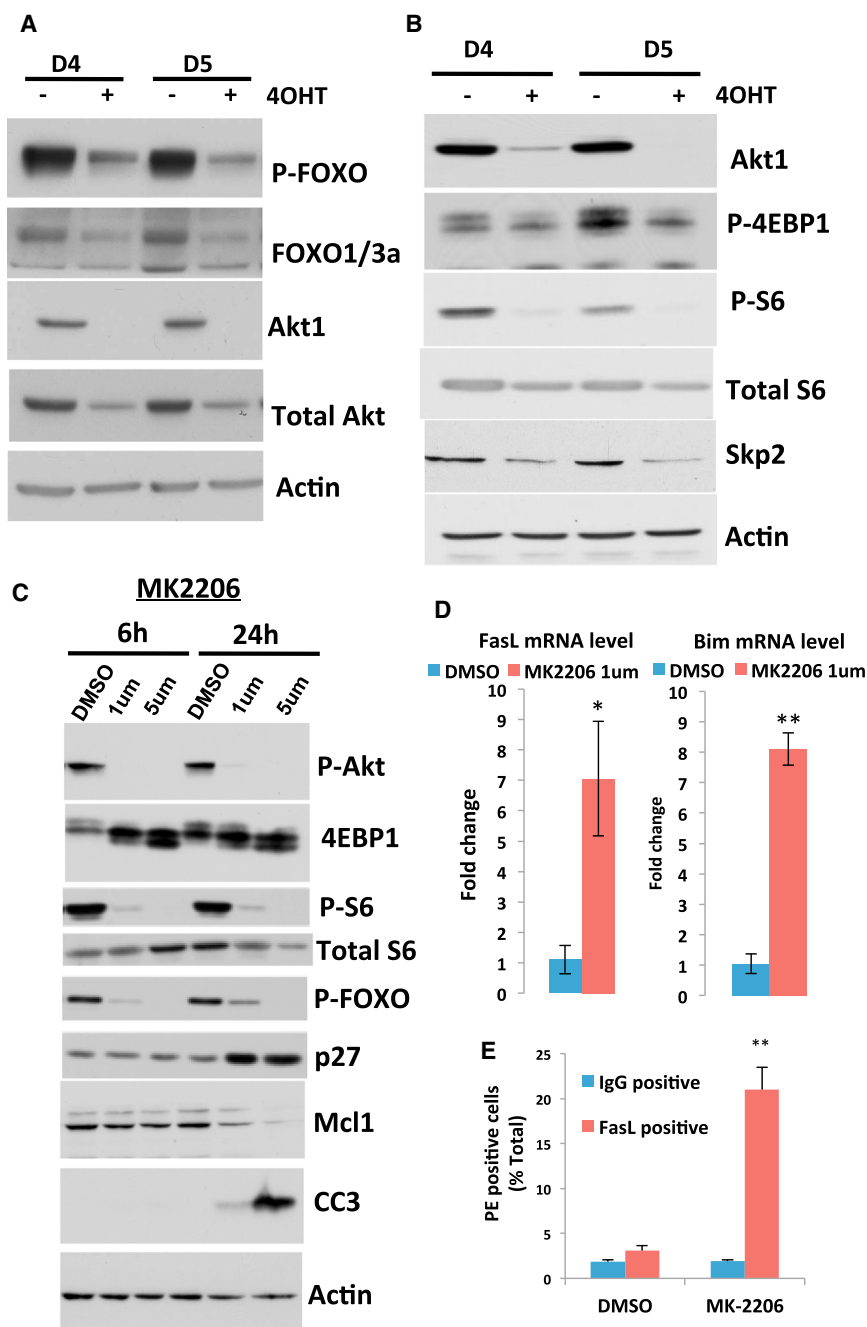
A

B

C

D

E

Figure 6. Activation of FOXO and Inhibition of mTORC1 and Skp2 Expression in Response to Akt1 Deletion or Akt Inhibition by MK2206

(A) FOXO phosphorylation at day 4 (D4) or day 5 (D5) after deletion of Akt1 by 4OHT in $p53^{-/-}$ Akt1f/f/R26CreERT2 lymphoma cells.

(B) Phosphorylation of 4EBP1, S6, and the expression of Skp2 at day 4 or 5 after deletion of Akt1 by 4OHT in $p53^{-/-}$Akt1f/f/R26CreERT2 lymphoma cells.

(C) Immunoblot showing FOXO phosphorylation, mTORC1 activity as measured by 4EBP1 mobility shift and S6 phosphorylation, p27 and MCL-1 protein levels, and CC3 after treatment of p53−53unoblot showing FO lymphoma cells with MK2206 at the indicated time points.

(D) Fas ligand (FasL) and Bim mRNA levels as measured by qRT-PCR after treatment of $p53^{-/-}$Akt1f/f/R26CreERT2 lymphoma cells with MK2206 for 18 hr. Data are presented as an average of three independent RNA preparations ±SEM. **$p < 0.001$, *$p < 0.05$.

(E) FACS analysis of FasL expression in cells treated with DMSO or 1 μM MK-2206 for 18 hr. Data are presented as an average of three independent experiments in triplicates ±SEM. **$p < 0.005$.

inhibited the induction of FAS and inhibited apoptosis induced by Akt inhibition (Figure 7F). All together, these results strongly indicate that the predominant initiator of apoptosis, induced by Akt inhibition in $p53^{-/-}$ thymic lymphoma, is FasL. Consistently, although we found that Bim mRNA levels are highly induced by the inhibition of Akt, we did not find a significant change in the major expressed isoform BIM-EL protein, and only small changes in the alternatively spliced isoform BIM-L (Figure S6B). Only in cells expressing activated Akt were the protein levels of the three BIM isoforms decreased.

Given that it was recently shown that D cyclins promote cell survival in hematopoietic cells by the inhibition of FasL and Fas receptor (Fas) expression (Choi et al., 2014), and since we did not find sig-

FasL in apoptosis, we monitored caspase-8 cleavage and found that it is increased in the control-treated cells but is inhibited in Skp2-overexpressing cells and mAkt-expressing cells (Figure 7D), to a similar extent as observed with caspase-3 cleavage (Figure 7B). Furthermore, inhibition of caspase-8 activity or FAS silencing inhibited cell death induced by Akt inhibition. We employed the caspase-8 inhibitor Z-LETD-FMK and showed that it markedly inhibited cell death induced by Akt inhibition (Figure 7E), without affecting the induction of FAS or FasL (Figure S5). Silencing FAS using small interfering RNA (siRNA) (Figure S6A)

nificant changes in BIM protein expression by Skp2 (Figure S6B), we analyzed cyclin D1 level in Skp2-overexpressing cells and found that it is significantly elevated (Figure 7G). The mechanism by which D cyclins inhibit FasL and Fas expression appears to be through the induction and activation of E2F1 (Choi et al., 2014). Therefore, the inhibition of p27 and the induction of cyclin D1 by Skp2 raise the possibility that Skp2 inhibits FasL mRNA via the phosphorylation of Rb and the activation of E2F. It is therefore expected that Skp2 overexpression would also reduce the expression of Fas. Indeed, we found that Skp2 overexpression

also markedly reduced the expression of Fas after Akt inhibition (Figure 7H). The results suggest a mechanism by which Skp2 antagonizes Akt-inhibition-mediated cell death in p53$^{-/-}$ thymic lymphoma (Figure 7I). Furthermore, the results uncovered a mechanism by which Skp2 could promote cell survival.

We clearly showed that the predominant mechanism, which selectively sensitizes p53$^{-/-}$ thymic lymphoma to cell death by Akt inhibition, is mediated by FAS and FASL. However, as we indicated above, it is possible that p53 deficiency in general sensitizes cells to cell death by Akt inhibition as was shown for the isogenic colorectal cancer cells (Figure 5F). Akt inhibition could induce cell death by multiple mechanisms, and the mechanism that we have uncovered in the thymic lymphoma cells may not apply to the colorectal cancer cell lines. We found that Akt inhibition, despite inducing higher levels of apoptosis in the p53-deficient human cancer cells, inhibits FOXO phosphorylation to a similar extent in p53-proficient and p53-deficient human cancer cells (Figure S7A). Unlike what was observed in the p53$^{-/-}$ thymic lymphoma cells, we did not find any significant and consistent changes in MCL-1 protein level upon Akt inhibition (Figure S7A). BIM and FAS protein levels were also did not consistently change upon Akt inhibition, although in the p53$^{-/-}$ HCC116 cells we found a modest elevation of FAS and BIM proteins (Figure S7A). We also did not find consistent changes in FasL expression. Only a modest increase of FasL mRNA was found in p53$^{-/-}$ SW40 and p53$^{-/-}$ DLD-1 cells when compared to the isogenic p53-proficient cells (Figure S7B). Nevertheless, caspase-3 cleavage was markedly induced in the p53-deficient cells when compared to the p53-proficient cells (Figure S7C). These results support the notion that regardless of the mechanism p53 deficiency sensitizes cells to cell death induced by the inhibition of Akt.

DISCUSSION

It is known that germline deletion of Akt1 is sufficient to inhibit tumor development in several mouse models of cancer, which are associated with Akt activation. However, these results do not address whether Akt1 inhibition is sufficient to inhibit or regress tumors after tumor onset. In addition, it is not clear whether Akt inhibitors could be therapeutic for cancers in which activation of Akt is not implicated. Akt hyperactivation is perhaps one of the most frequently occurring events in human cancers, and therefore these cancers are rationale candidates for Akt inhibition therapy, but it is not known whether Akt inhibition is valuable in cancers, which do not display activation of Akt. Finally, it is not clear what are the potential mechanisms that could be developed by cancer cells to resist Akt inhibition therapy. In the current studies, we addressed these three questions and showed that systemic deletion of Akt1 in mice is therapeutic for p53$^{-/-}$ thymic lymphoma that does not display Akt activation.

The effect of systemic Akt1 deletion on the lifespan extension of p53$^{-/-}$ tumor bearing mice is profound but is not sufficient to reach normal lifespan. Although the MRI studies showed that the systemic Akt1 deletion either halt the growth or regress the p53$^{-/-}$ thymic lymphoma, it is possible that other tumors developed in p53$^{-/-}$ mice caused the eventual mortality. Usually p53$^{-/-}$ mice also develop high incidence of sarcoma, but we

have not detected sarcomas within the time frame that the mice were treated possibly because of the C57Bl6 background of the mice used in our studies. We did observe, however, sarcomas in older mice, which could contribute to the mortality even after Akt1 deletion. Alternatively or additionally, thymic lymphoma cells that escaped the deletion could contribute to the mortality. Finally, the possibility that p53$^{-/-}$ mice have a shorter lifespan regardless of tumor development cannot be completely excluded. Future studies are required to determine whether other types of tumors, which are driven by p53 deletion or by mutated p53, such as sarcoma, could be also targeted by Akt1 inhibition.

The germline deletion of Akt1 in the mouse has only minor consequences, and, in fact, Akt1$^{-/-}$ mice live longer than wild-type mice (Chen et al., 2006). The systemic deletion of Akt1 in adult mice does not induce any overt phenotype, and therefore our results strongly suggest that Akt1 specific inhibition could be well tolerated and it is an efficient cancer therapy strategy even if the cancer cells do not display Akt activation. The results provide a paradigm that might expand the spectrum of cancer types that could be candidates for Akt inhibition therapy. Restoration of p53 had been suggested as a strategy for cancer therapy (Martins et al., 2006; Ventura et al., 2007; Wang et al., 2011). However, restoration of p53 is difficult to achieve, and thus our results suggest a feasible strategy for therapy of cancers that are driven by the loss of p53. Moreover, our results could be extended to human cancer cells deficient for p53 as we found that p53 deficiency sensitizes human cancer cells to cell death induced by Akt inhibition.

As expected, we found a decrease in FOXO phosphorylation and mTORC1 activity after Akt1 deletion or Akt inhibition in p53$^{-/-}$ thymic lymphoma. The activation of FOXO resulted an increase in its pro-apoptotic transcriptional targets Bim and FasL. The inhibition of Akt also reduced the protein levels of Mcl-1. Akt inhibition could reduce the steady-state level of Mcl-1 through inhibition of its mRNA translation, which is highly dependent on eIF4E (Wendel et al., 2007) or through its destabilization by GSK3, which is phosphorylated and inactivated by Akt (Maurer et al., 2006). The combination of the elevated Bim and FasL with the reduced Mcl-1 level could explain the cell death induced by Akt inhibition in p53$^{-/-}$ thymic lymphoma. However, our results strongly indicated that the predominant cause of cell death is FasL mediated. Skp2 expression was also reduced after Akt inhibition. The reduced Skp2 expression is likely due to reduced Skp2 mRNA translation since it is highly dependent on Akt and on its downstream effectors mTORC1 and eIF4E, as we had shown previously (Nogueira et al., 2012). However, we observed that in the thymic lymphoma cells exogenous Skp2 protein is also reduced upon Akt inhibition, which cannot be explained only by mRNA translation. Other results showed that Skp2 is directly phosphorylated by Akt, which in turn maintains its cytoplasmic localization and stability (Gao et al., 2009; Lin et al., 2009), but these results were challenged by others (Bashir et al., 2010). We expected that ectopic Skp2 expression would override the inhibition of cell proliferation induced by Akt inhibition in p53$^{-/-}$ thymic lymphoma. Surprisingly, we found that high level of ectopic Skp2 expression in p53$^{-/-}$ thymic lymphoma exerts resistance to cell death induced by Akt inhibition. Indeed, previous reports documented that silencing of Skp2 in a
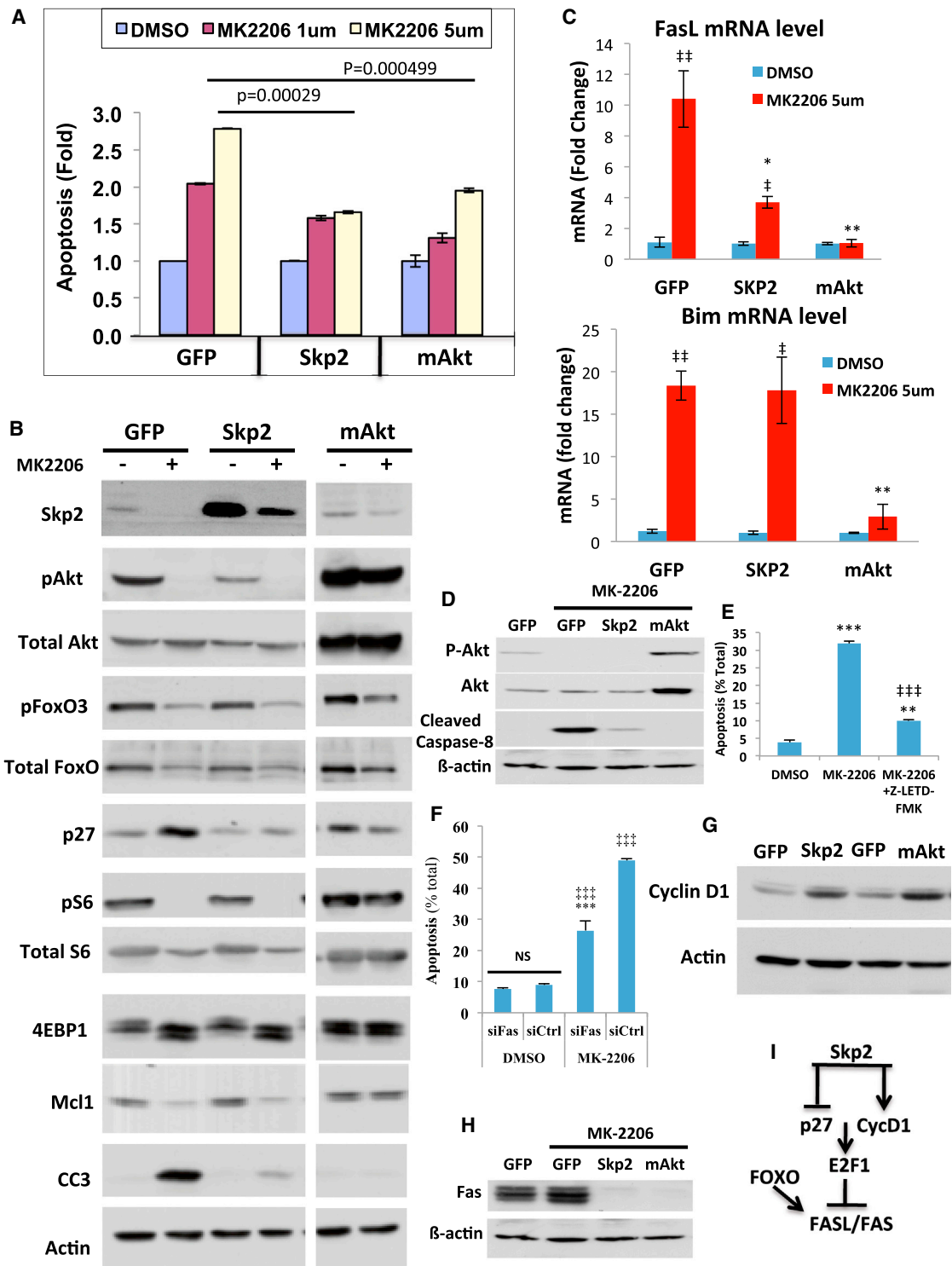
Figure 7. Ectopic Expression of Skp2 or mAkt in p53$^{-/-}$ Thymic Lymphoma Cells Exerts Resistance to MK2206

(A) Induction of cell death of control GFP-expressing cells, Skp2, or mAkt-expressing cells after treatment with MK2206 for 18 hr. Data are presented as an average of three independent experiments ±SEM.

(B) Immunoblot showing the phosphorylation of Skp2 expression, FOXO3 phosphorylation, p27 expression, mTORC1 activity as measured by S6 phosphory-lation and 4EBP1 mobility shift, Mcl-1 and cleaved caspase-3 (CC3) expression in GFP-, Skp2-, or mAkt-expressing cells following treatment with MK2206 for 18 hr.

(legend continued on next page)

variety of cancer cell lines elicits cell death (Jiang et al., 2005; Lee and McCormick, 2005). In particular, pRb-deficient cancer cells are sensitized to apoptosis induced by Skp2 silencing with no effect on proliferation. These results suggest that Skp2 has a role in cell survival, which is independent from its role in cell proliferation (Wang et al., 2010a). Interestingly, high Skp2 expression was implicated in T cell lymphoma in mice and in high-grade lymphomas in human (Latres et al., 2001). Expression of Skp2 in T cells in mice profoundly increased the incidence of T cell lymphoma induced by oncogenic N-Ras (Latres et al., 2001), and mice lacking CREB binding protein (CBP) develop T cell lymphomagenesis with concomitant elevated Skp2 expression (Kang-Decker et al., 2004). We found that Skp2 decreased the induction of FasL and Fas by MK2206 treatment. Our results showed that the predominant mechanism by which Akt inhibition promotes cell death in $p53^{-/-}$ thymic lymphoma is via FASL and FAS. Both FAS silencing and caspase-8 inhibition inhibited the cell death induced by Akt inhibition. Consistently, it was reported that the deficiency of FasL accelerates development of tumors in $p53^{+/-}$ mice (Embree-Ku and Boekelheide, 2002). Skp2 overexpression negates apoptosis induced by Akt inhibition by prohibiting the induction of FasL and by reducing the expression of FAS. As was recently reported, D cyclins promote hematopoietic cell survival through inhibition of FasL and Fas expression via E2F1 (Choi et al., 2014). Since we found that Skp2 overexpression induced cyclin D1 expression and inhibits p27 expression, which eventually could activate E2F1, we proposed a mechanism by which Skp2 negates Fas-mediated apoptosis induced by FOXO (Figure 7I). Thus, Skp2 exerts resistance to Akt inhibition and could override both inhibition of mTORC1 and activation of FOXO and could predict resistance to Akt inhibition therapy. Consistently, it was reported that resistance to rapamycin is linked to high Skp2 expression (Totary-Jain et al., 2012). Although Skp2 had been previously implicated in inhibition of apoptosis, the mechanism by which it exerts resistance to apoptosis is unclear. Our results provided a mechanism by which Skp2 could promote resistance to FAS-mediated cell death. The hypersensitivity to Akt inhibition exerted by p53 deficiency, although it was demonstrated only in one particular mouse model of $p53^{-/-}$ thymic lymphoma, could be potentially generalized to other cell types in future work. However, the mechanism by which Akt inhibition sensitizes $p53^{-/-}$ human epithelial cancer cells to cell death appears to be different from the mechanism uncovered in the thymic lymphoma cells, and could be dependent on the cells' milieu.

In summary, the results presented in this study provided a concept showing that systemic Akt1 inhibition alone is sufficient to selectively regress and inhibit tumor progression even in the absence of Akt activation in this tumor. It was also emerged from these studies that deregulation of Skp2 expression is a potential mechanism for resistance to Akt inhibition.

EXPERIMENTAL PROCEDURES

Reagent and Antibodies
RPMI-1640 was purchased from Gibco and HyClone. Fetal bovine serum (FBS) was purchased from Gemini. MK2206 was purchased from Selleckchem and ChemieTek. Captisol was purchased from Captisol, a Ligand Company. Antibodies against phospho-Akt (S473), Akt, phospho-S6, S6, phospho-FoxO3a, phospho-4EBP1, 4EBP1, cleaved caspase-3, and Gadph from Cell Signaling Technology. Antibodies to actin from Sigma; Skp2, BIM, Cyclin D1, and cleaved caspase-8 antibodies are from Cell Signal and Abcam; FoxO3 antibodies are from Upstate, p27 antibodies are from BD Biosciences, and Mcl-1 and FAS antibodies are from Santa Cruz Biotechnology. BrdU was purchased from Dako, and fluorescein mouse second antibody was from Vector Laboratories.

Mice
C57Bl/6 $Akt1^{f/f}$ mice were generated by Ozgene (Australia) as described in Figure S1. The Rosa26CreERT2 strain (Strain 01XAB) was obtained from Tyler Jacks' laboratory (Massachusetts Institute of Technology). The C57Bl/6 $p53^{+/-}$ strain was obtained from the Jackson Laboratory. The double-compound $R26CreERT2; p53^{-/-}$ mice and triple-compound $R26CreERT2; Akt1^{f/f}; p53^{-/-}$ mice were generated by mating the three individual strains. NU/NU nude mice were obtained from Charles River Laboratories.

Genotyping
DNA samples from tail of mice or thymic lymphoma cells were analyzed by PCR using primer sets that can detect wild-type, deleted p53, Akt1 floxed, and Rosa26CreERT2 alleles. Primers sequences are as follows: for Akt1 floxed allele, 5′-TCACAGAGATCCACCTGTGC-3′(A1-3loxp), 5′-GGGCCTCCATACA CTCAAGA-3′(A1-i4-5R); p53, ACAGCGTGGTGGTACCTTAT, TCCTCGTGCT TTACGGTATC, TATACTCAGAGCCGGCCT; CreERT2, CCTGATCCTGGCAA TTTCG (CRE-R1), GGAGCGGGAGAAATGGATATG (CRE-wtR), and AAAGTC GCTCTGAGTTGTTAT (CRE-uniF).

Isolation of Primary Thymocytes and Establishment of p53-Null Thymic Lymphoma Cell Lines
Primary thymocytes were isolated by gently pressing the thymus between two sterile glass slides and then washed with PBS and resuspended in RPMI medium supplemented with 10% fetal bovine serum (FBS) and penicillin-streptomycin. For the generation of thymic lymphoma cell lines, tumors were excised and minced. Thymic lymphoma cells grew in suspension, and they were maintained in RPMI medium supplemented with 10% fetal bovine serum (FBS) and penicillin-streptomycin.

Isogenic Colorectal Cell Lines
The cell lines HCT116, HCT116-$p53^{-/-}$, DLD1, DLD-$p53^{-/-}$, SW48, and SW48-$p53^{-/-}$ were previously described (Sur et al., 2009) and were kindly provided by Dr. Bert Vogelstein. Cells were grown in DMEM medium

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(C) Expression of FasL and Bim as measured by RT-PCR in GFP-, Skp2-, or mAkt-expressing cells following treatment with MK2206 for 18 hr. Data are presented as an average of three independent RNA preparations ±SEM. **p < 0.01, *p < 0.05 versus GFP; ‡‡p < 0.01, ‡p < 0.05 versus DMSO in each group.

(D) Immunoblot showing the level of cleaved caspase-8 in GFP-, Skp2-, or mAkt-expressing cells following treatment with 1 μM MK2206 for 18 hr.

(E) Induction of cell death of $p53^{-/-}$ thymic lymphoma cells after treatment with 1 μM MK2206 for 18 hr in presence or absence of cell permeable caspase-8 inhibitor (Z-LETD-FMK). Data are presented as an average of three independent experiments in triplicates ±SEM. ***p < 0.0001 and **p < 0.005 compared to DMSO. ‡‡‡p < 0.0001 compared to MK2206.

(F) Cells were treated with DMSO or MK2206 for 18 hr after knockdown of Fas using a pool of four siRNAs. Following treatment, cell death was assessed by flow (PI staining). Experiments were done twice in triplicate. Statistical analysis: ***p < 0.0001 compared to control siRNA. ‡‡‡p < 0.0001 compared to DMSO.

(G) Immunoblot showing Cyclin D1 expression in GFP-, Skp2-, or mAkt-expressing cells.

(H) Fas expression in GFP-, Skp2-, or mAkt-expressing cells.

(I) Illustration depicting the mechanism by which Akt inhibition promotes cells death in $p53^{-/-}$ thymic lymphoma.

supplemented with 10% FBS. For treatment with MK2206, cells (300,000) were plated on 6-cm plates and 12 hr later were exposed to either MK2206 or DMSO. Apoptosis was quantified with DAPI attaining as previously described (Kennedy et al., 1997).

qPCR

Total RNA was extracted from cells at the indicated treatment and time point by using TRIzol reagent (Invitrogen). First-strand cDNA was produced with SuperScript III reverse transcriptase (Invitrogen). Quantitative real-time PCR was performed with iQ SYBR green supermix (BIO-RAD) with the iQ5 real-time PCR detecting system. Relative levels of mRNA were compared by using actin or cyclophilin A primers to normalize for total RNA input. The reaction conditions were as follows: 5 min of denaturation at 95°C, followed by 45 cycles of denaturation for 15 s at 95°C, and annealing and elongation for 1 min at 55°C. The mouse primer sequences are as follows: Bim, 5'-CGA CAGTCTCAGGAGGAACC-3' (forward) and 5'-CCTTCTCCATACCAGACG GA-3' (reverse); FasL, 5'-TCCGTGAGTTCACCAACCAAA-3' (forward) and 5'-GGGGGTTCCCTGTTAAATGGG-3' (reverse); Cycophilin A, 5'-TTCACAAA CCACAATGGCACAGGG-3' (forward) and 5'-TGCCGTCCAGCCAATCTGT CTTAT-3' (reverse); Actin, 5'-GGCCGAGGAGCAAGAATGG-3' (forward) and 5'-CATGCACTCTGCGATACGCT-3' (reverse).

Western Blot Analysis

Cells were collected and lysed in lysis buffer (1% Triton X-100, 20 mM HEPES, 150 mM NaCl, 1 mM EDTA, 1 mM EGTA) containing phosphatase inhibitors (5 mM IAA, 20 nM OA, 10 mM sodium pyrophosphate, 20 mM β-glycero-glycerophosphate,100 mM NaF) and protease inhibitor cocktail (Roche Applied Science). Solubilized proteins were collected by centrifugation and quantified using a protein assay reagent (Bio-Rad). Equal amounts of protein of each sample were resolved by electrophoresis in 8%, 10%, or 12% gel and transferred to polyvinylidene difluoride membranes (Bio-Rad).

Retroviral Plasmids

The retroviral vector p-MSCV-IRES-GFP was obtained from Addgene (Addgene plasmid 20672). Myristoylated Akt1 (mAkt) was previously described (Skeen et al., 2006) and was inserted in the Sal1 and Xho1 of p-MSCV-IRES-GFP. The retroviral vector p-MSCV-Skp2-IRES-GFP was kindly provided by Michael Deininger and was previously described (Agarwal et al., 2008). The retroviral vector p-MSCV-p53-IRES-GFP was kindly provided by Dean Felsher and was previously described (Giuriato et al., 2006).

Retrovirus Transfection and Infection

Retroviruses were produced in phoenix-ecotropic cells by lipofectamine 2000 transfection. Thymic lymphoma cells were infected by using MagnetoFection. Before infection, viruses were mixed with ViroMag R/L and incubated at room temperature for 15 min. Cells were mixed with CombiMag and then plated in 12-well plates and incubated on a magnetic plate for 20 min. After incubation, ViroMag R/L/viruses mixture were added to the CombiMag/cells mixture and incubated for 30–60 min following manufacturer's instructions. Seven days after infection, GFP-positive cells were sorted using MoFlo Legacy cell sorter to establish stably expressing cell lines.

Transfection with siRNA

Cells were transfected using MagnetoFection. Two siRNAs pool were used: non-targeting control siRNA (ON-TARGETplus non-targeting siRNA SMARTpool, Dharmacon) or a siRNA pool consisting of four siRNAs specific for mouse Fas (ON-TARGETplus Fas siRNA SMARTpool, Dharmacon). Before infection, siRNA were mixed with SilentMag and incubated at room temperature for 15 min. In the meantime, cells were mixed with CombiMag, plated in a 12-well plate and incubated on the magnetic plate for 20 min. After incubation, siRNA/SilentMag mixture was added to the CombiMag/cells mixture on the magnetic plate for 30 min as recommended by manufacturer's instructions. Three days post transfection cells were split for experiment and treated with DMSO or MK2206 for 18 hr. Flow cytometry was then performed to estimate cell death (PI staining), and samples were collected for immunoblotting.

Flow Cytometry, Cell-Cycle Analysis, BrdU Incorporation, and Cell Sorting

Cells were incubated with bromodeoxyuridine (BrdU) (3 μg/ml) for 2 hr before the indicated time point for collection. Cells were resuspended in 1 ml PBS after centrifugation and fixed by adding 3 ml of 100% ethanol dropwise and stored at 4°C. The day before analysis, cells were pelleted by centrifugation at 2,000 rpm for 5 min. Then cells were solubilized in 2N-HCl with 0.5% Triton-X followed by neutralization with 0.1 M sodium borate (pH 8.5). Cells were then incubated with anti-BrdU antibody (Dako) overnight at 4°C. The next day, cells were washed twice by washing buffer (PBS + 0.5% BSA) and incubated with fluorescein anti-mouse secondary antibody (Vector Labs). Before analysis, the cells were stained with propidium iodide containing RNase and Triton X-100. Flow cytometry was performed using Beckman Coulter EPICS Elite ES (Beckman) and analyzed using Cell lab Quanta SC. Cell sorting was performed using MoFlo Legacy cell sorter.

Assessment of FasL Cell-Surface Expression

Cells were incubated with DMSO or MK-2206 (1μM) for 18 hr before collection to assess FasL surface expression by FACS. For FasL staining, cell pellets were resuspended in cell staining buffer (BioLegend) and incubated with 1 μg/3 × 10^6 cells PE anti-mouse FasL antibody (BioLegend) for 1 hr at 4°C in the dark. The cells were then washed twice in cell staining buffer and counterstained with 7-AAD (500 ng/3 × 10^6 cells) for another 15 min at 4°C in the dark to assess viability. As a negative control, cells were stained with PE IgG isotype control antibody. Flow cytometry was then performed using Cell lab Quanta SC, and both FasL expression (PE positive cells) and apoptosis (7-AAD positive cells) were quantified.

Caspase-8 Inhibition

Cells were incubated with DMSO or MK-2206 (1 μM) in absence or presence of 20 μM caspase-8 inhibitor (Z-LETD-FMK, Bio-Rad) for 18 hr before collection. The cells were then washed twice in PBS/5 mM EDTA and stained with 7-AAD (500 ng/3 × 10^6 cells) for 15 min at 4°C in the dark to assess viability. Flow cytometry was then performed using Cell lab Quanta SC, and apoptosis (7-AAD positive cells) was quantified.

Xenograft Assay

Cells were counted and suspended in cold PBS. Cells (1.5 × 10^6) were injected subcutaneously into the rear flank of the nude mice. After palpable tumor formation, mice were randomized into two groups. Either corn oil or tamoxifen (1 mg/100 μl/per injection) was injected into the nude mice intraperitoneally every other day for 2 weeks. MK2206 was dissolved in 30% captisol per manufacturer's instructions and delivered by oral gavage. Tumor sizes were measured with a caliper and calculated by length × height × width × 0.5.

MRI and Tumor Volume Analysis

All animals were sequentially imaged using an Agilent 9.4T preclinical MRI to screen for tumor growth and treatment response to tamoxifen. T1-weighted sequences were obtained in axial and coronal planes, using the following parameters: T1-weighted: repetition time (TR)/echo time (TE) = 900/14 ms, matrix size 192 × 192, 1-mm slice thickness, 20 sequences, six repetitions. Tumor volume measurements were performed using T1-weighted coronal and axial image stacks. The tumor volume was calculated by ImageJ measure stacks to obtain the tumor volume in cubic millimeters. For each time point, all measurements from the available sequences were used to calculate a mean volume ±SEM in order to limit errors due to manual process.

Statistical Analysis

Statistical analysis was done with unpaired Student's t tests. Survival curves were analyzed by log-rank test (Mantel-Cox), and data are expressed as SEM as indicated in the figure legends. Unless otherwise indicated, all the experiments were performed for at least three times.

SUPPLEMENTAL INFORMATION

Supplemental Information includes seven figures and can be found with this article online at http://dx.doi.org/10.1016/j.celrep.2015.06.057.

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