Reciprocal Feedback Regulation of PI3K and Androgen Receptor Signaling in PTEN-Deficient Prostate Cancer

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SUMMARY

Prostate cancer is characterized by its dependence on androgen receptor (AR) and frequent activation of PI3K signaling. We find that AR transcriptional output is decreased in human and murine tumors with PTEN deletion and that PI3K pathway inhibition activates AR signaling by relieving feedback inhibition of HER kinases. Similarly, AR inhibition activates AKT signaling by reducing levels of the AKT phosphatase PHLPP. Thus, these two oncogenic pathways cross-regulate each other by reciprocal feedback. Inhibition of one activates the other, thereby maintaining tumor cell survival. However, combined pharmacologic inhibition of PI3K and AR signaling caused near-complete prostate cancer regressions in a Pten-deficient murine prostate cancer model and in human prostate cancer xenografts, indicating that both pathways coordinately support survival.

INTRODUCTION

Prostate cancer is the most common malignancy diagnosed in males and the second most common cause of male cancer deaths (American Cancer Society, 2010). Despite advances made in the early detection and treatment of localized prostate cancer, the American Cancer Society estimates that 32,050 men will have died of metastatic disease in 2010. Androgen-deprivation therapy remains the standard treatment of metastatic prostate cancer; however, progression to castrate resistance disease occurs in the majority of patients (Rini and Small, 2002; Singh et al., 2010). Following the emergence of castrate-resistant prostate cancer, docetaxel chemotherapy has been shown to be therapeutically efficacious; however, the median increase in survival was only 4 months (Petrylak et al., 2004). Thus, there is a significant need for improvements in therapy for prostate cancer.

The PI3K pathway plays a central role in tumorigenesis across a variety of malignancies (Courtney et al., 2010; Liu et al., 2009). Prostate cancers are associated with genetic alterations involving the PI3K and androgen receptor (AR) pathways, both of which mediate survival signals in prostate cancer. Roughly 40% of primary and 70% of metastatic prostate cancers have genomic alterations in the PI3K-signaling pathway, mostly through loss of PTEN (El Sheikh et al., 2008; Reid et al., 2010; Taylor et al., 2010). Preclinical studies of mice with conditional,
prostate-specific Pten deletion and of cell lines with stable silencing of Pten by RNA interference have established that loss of PTEN promotes resistance to castration (Gao et al., 2006; Jiao et al., 2007). However, this effect of PTEN loss is not absolute because certain prostate cancer xenograft models with PTEN loss (LNCaP, LAPC9) remain at least partially sensitive to castration (Craft et al., 1999). Furthermore, the high clinical response rate to castration therapy indicates that at least some PTEN-deficient tumors retain some degree of sensitivity.

The critical role of PTEN in regulating flux through the PI3K-signaling pathway raises the possibility that PI3K pathway inhibitors might be effective in PTEN-deficient prostate cancer. Indeed, genetic loss of either mTOR or AKT1 is sufficient to significantly reduce the initiation of prostate cancer in the conditional Pten model (Chen et al., 2006; Guertin et al., 2009; Nardella et al., 2009). The mTORC1 inhibitor rapamycin has been shown to revert early PIN lesions in young mAKT mice; however, results in Pten prostate conditional null mouse models have been modest (Majumder et al., 2004; Zhang et al., 2009). Furthermore, clinical trials of rapamycin analogs in castration-resistant prostate cancer have failed to demonstrate clinical activity (George et al., 2008, J. Clin. Oncol., abstract). One potential liability of mTORC1 inhibition is disruption of a negative feedback loop, resulting in hyperactivation of AKT and MAPK that can promote cell survival independent of mTORC1, thereby limiting therapeutic efficacy (Carracedo et al., 2008; O'Reilly et al., 2006; Sarbassov et al., 2005).

The availability of a number of PI3K pathway inhibitors in clinical development targeting various critical components of the pathway (PI3K, AKT, mTORC1/2) allows this issue to be readdressed (Serra et al., 2008). The goal of our study was to evaluate the therapeutic efficacy of PI3K pathway inhibition in preclinical models of prostate cancer and to define the molecular mechanism of PI3K and AR feedback regulation. Through this work we propose combination therapy based on targeting compensatory survival pathways associated with relief of feedback inhibition observed following PI3K or AR inhibition.

RESULTS

Inhibition of the PI3K Pathway Causes Growth Arrest but Not Significant Tumor Regression in Pten-Negative Prostate Cancers

We evaluated the therapeutic efficacy of PI3K pathway inhibition in mice with established prostate cancers caused by either conditional deletion of Pten (Pten<sup>lox/lox</sup> PB-Cre, hereafter called Pten<sup>lox/lox</sup>·Cre) or transgenic expression of MYC (PB-MYC) using BEZ235, a dual PI3K and mTORC1/2 inhibitor (Serra et al., 2008). PB-MYC mice were chosen because MYC amplification or overexpression is also commonly found in human tumors. This model likely represents a subset of human prostate cancer distinct from that driven by PTEN loss. PI3K/mTOR inhibition was confirmed in the Pten<sup>lox/lox</sup> mice using pAKT and pS6 and in the PB-MYC mice using pS6 (Figure 1A; see Figure S1A available online). Cell proliferation as measured by Ki67 staining was significantly reduced in the Pten<sup>lox/lox</sup> mice (proportion of positive cells: 0.34 versus 0.15, for vehicle control and BEZ235, respectively; p < 0.01), but not in PB-MYC mice (Figures 1B and 1C). However, there was minimal reduction in prostate cancer tumor volume as measured by MRI (mean volume reduction of 14.75%) and no obvious effect on tumor histology (Figures 1B, 1D, and 1E). PB-MYC prostate cancers showed no radiographic or histologic response (Figures 1B–1E). In summary, BEZ235 has modest, primarily cytostatic, activity in Pten<sup>lox/lox</sup> mice, but no activity in PB-MYC mice, consistent with earlier in vitro studies in breast cancer cell lines (Brachmann et al., 2009).

Inhibition of the PI3K Pathway Activates the AR Pathway in PTEN-Negative Prostate Cancers

Given the critical role of AR in prostate cancer initiation and progression, we hypothesized that sustained AR activity might explain the persistent survival of Pten null prostate cells in Pten<sup>lox/lox</sup> mice treated with BEZ235. To our surprise, we found that Pten<sup>lox/lox</sup> mice had reduced AR protein levels compared to their Pten wild-type littermates. Treatment of Pten<sup>lox/lox</sup> mice with BEZ235 partially rescued AR protein levels, indicating that increased PI3K/mTOR activity likely explains the decrease in AR levels (Figure 2A). Similar effects of PI3K/mTOR inhibition (BEZ235) or mTORC1 (RAD001) inhibition on AR protein levels were observed in the PTEN-deficient human prostate cancer cell line LNCaP (Figure 2B). As expected from earlier studies with rapamycin (Carracedo et al., 2008), p-ERK levels were increased following treatment with either BEZ235 or RAD001 (Figure 2B). Thus, PI3K pathway inhibition in PTEN-deficient prostate cancer resulted in the activation of two critical cell survival pathways (AR and MAPK).

We next evaluated whether the increase in AR protein levels seen with PI3K pathway inhibition resulted in increased AR target gene activity. Indeed, mRNA levels of three canonical AR target genes, Pbsn, Nkx3.1, and Psca, were increased by short-term treatment of Pten<sup>lox/lox</sup> mice with BEZ235 (Figure 2C). Similarly, the activity of an androgen-responsive reporter gene was increased in LNCaP cells exposed to BEZ235 or RAD001, consistent with other reports using rapamycin (Figure 2D) (Lin et al., 2003; Wang et al., 2008). Increased androgen-responsive reporter gene activity was also observed following treatment with an allosteric, highly specific, inhibitor of AKT1 and AKT2, providing further evidence that these pharmacologic effects are due to PI3K pathway blockade (Figure 2D). Collectively, the data from these PTEN-deficient murine and human models indicate that PI3K pathway inhibitors can activate AR target gene expression.

HER family receptor tyrosine kinases and the insulin-like growth factor receptor (IGF1R) are feedback inhibited by the PI3K pathway and reactivated in breast tumor cell lines and xenograft models exposed to AKT inhibitors (Chandarlapaty et al., 2011). Consistent with these data, others have found that inhibition of PI3K pathway in breast cancer cell lines results in upregulation of HER3 (Yao et al., 2009). Therefore, we considered that similar effects may be seen in PTEN-deficient prostate cancers. Indeed, the level of HER3 was increased in LNCaP cells and in prostates of Pten<sup>lox/lox</sup> mice after exposure to BEZ235 (Figure 2E). To address potential off-target effects associated with pharmacologic inhibition of PI3K pathway kinases, we targeted AKT1/2 using siRNA. AKT1/2 knockdown led to upregulation of both HER3 and AR, as well as pERK similar to that...
observed with BEZ235, albeit to more modest levels (Figure 2F).

We and others previously demonstrated that HER2/HER3 promotes AR activity and stability by an AKT-independent mechanism by enhancing both AR stability and transcriptional activity, similar to the effects observed here with BEZ235 (Mahajan et al., 2007; Mellinghoff et al., 2004; Yeh et al., 1999). Therefore, we postulated that the increase in HER3 expression induced by PI3K pathway inhibition might explain the increase in AR transcriptional output. Consistent with this hypothesis, cotreatment with the HER family kinase inhibitor PKI166 (EGFR and HER2 inhibitor) abolished the upregulation of AR activity observed with BEZ235, RAD001, or AKT1/2 inhibitor (Figure 2G).

To address the potential role of ERK activation in the upregulation of AR activity, we examined the effects of MEK inhibition using the MEK inhibitor PD0325901. In contrast to the stimulatory
Figure 2. Inhibition of the PI3K Pathway Restores, in Part, Androgen-Responsive Signaling in PTEN-Loss Prostate Cancers

(A) AR protein levels by western blot analysis in wild-type, Pten$^{lox/lox}$, and Pten$^{lox/lox}$ mice treated with BEZ235. Ventral prostates from individual 8-week-old mice were analyzed.

(B) Western blot analysis of AR, pAKT, pS6, and pERK in the LNCaP cell line treated for 24 hr with DMSO, BEZ235 (500 nM), and RAD001 (100 nM).

(C) Expression analysis (qRT-PCR) of the androgen-regulated genes Pbsn, Nkx3.1, and Psca in wild-type, Pten$^{lox/lox}$, and Pten$^{lox/lox}$ mice treated with BEZ235, normalized to Hprt and wild-type intact mice. Ventral prostates from individual 8-week-old mice were analyzed, and mean fold change ±SD is reported.

(D) Luciferase mean fold change ±SD measurements in the LNCaP AR-ARE-Luciferase cell line treated for 24 hr with DMSO, BEZ235 (500 nM), RAD001 (100 nM), and AKT1/2 inhibitor (1 μM) normalized to DMSO.

(E) Western blot analysis of HER2, HER3, AR, and PSA in the LNCaP cell line treated with DMSO and BEZ235 (500 nM) for 24 hr and Pten$^{lox/lox}$ mice treated with BEZ235.

(F) Western blot analysis of HER3, AR, pAKT, and pERK in LNCaP cells treated with BEZ235 or siRNA AKT1/2 for 24 hr.

(G) Luciferase mean fold change ±SD measurements in the LNCaP AR-ARE-Luciferase cell line treated for 24 hr with DMSO, BEZ235 (500 nM), RAD001 (100 nM), with or without PKI166 (5 μM), or PD0325901 (1 μM) normalized to DMSO.
effects of BEZ235 and RAD001 on AR signaling, PD0325901 did not augment AR signaling. Furthermore, PD0325901 did not reverse AR activation induced by BEZ235 or RAD001, despite the fact that ERK is activated by both drugs (Figure 2G). Thus, inhibition of the PI3K pathway upregulates AR target gene expression in a HER kinase-dependent manner independent of MEK.

PTEN Loss Is Associated with Repression of Androgen-Responsive Genes

Having demonstrated that inhibition of the PI3K pathway results in increased AR activity in two prostate cancer models, we explored the relevance of this finding in human prostate cancer specimens. Because clinical trials of PI3K pathway inhibitors in prostate cancer are still in early stages, we asked the reciprocal question of whether PI3K activation caused by PTEN loss impairs AR activity in primary human prostate tumors. One hundred and six tumors from a previously reported Memorial Sloan-Kettering Cancer Center (MSKCC) data set were designated PTEN loss or PTEN normal based on PTEN copy number and PTEN mRNA expression level. These PTEN status assignments were validated by gene set enrichment analysis (GSEA) showing concordance with a transcriptome-based signature of PTEN loss developed independently from breast cancer specimens (Figure S2A) (Saal et al., 2007). We then analyzed AR pathway activation by PTEN status using a previously reported mRNA signature of AR target genes (Hieronymus et al., 2006). AR activity was significantly repressed in PTEN-loss prostate tumors (Student’s t test on summed z score, p < 0.001; Figure 3A). Consistent with this finding, GSEA of a murine androgen-responsive gene set in Pten-loss prostate cancers compared to an intact wild-type mouse. See also Figure S2 and Tables S1 and S2.

Figure 3. PTEN-Loss Prostate Cancer Is Associated with Reduced Androgen-Responsive Gene Signaling

(A) Mean summed z score ±SD for an androgen-responsive gene signature in human PTEN-loss and PTEN-normal prostate cancer showing that PTEN-loss tumors have a reduced androgen-responsive gene signature with corresponding heat map. Significance was determined by the Student’s t test. (B and C) GSEA showing the (B) Hieronymus et al. (2006) and (C) Nelson et al. (2002) androgen-responsive gene signatures revealed that androgen-responsive gene signaling is reduced in PTEN-loss human prostate cancers. (D) Mean ERBB2 expression levels ±SD in PTEN-loss and PTEN-normal prostate cancer. Significance was determined by the Student’s t test. (E) GSEA of a murine androgen-responsive gene set in Pten-loss prostate cancers compared to an intact wild-type mouse. See also Figure S2 and Tables S1 and S2.
revealed that the same androgen-regulated gene set was significantly repressed in the PTEN-loss cancers (FDR = 0.01, Figure 3B) (Hieronymus et al., 2006). This association was also observed with two other independently derived AR target gene sets (FDR = 0.01 [Nelson et al., 2002], Figure 3C; FDR < 0.001 [Creighton, 2008], data not shown).

Our observation that PI3K inhibition leads to increased HER3 levels in Ptenlox/lox mice and in LNCaP cells raises the possibility that human tumors with PTEN loss might have decreased HER2/3 activity. We did not observe significant differences in HER3 mRNA levels, but HER2 (ERBB2) expression was significantly (p = 0.006) reduced in PTEN-loss prostate cancers (Figure 3D). Furthermore, HER2 expression was significantly (p < 0.003) correlated with AR target gene signature output (Figure S2B).

Because other genomic alterations may impact the interpretation of the human tumor studies, we examined AR activity in primary prostate tissue harvested from 8-week-old Ptenlox/lox mice (with or with Pten deletion) before the onset of prostate cancer. To define a murine AR gene signature, we first compared transcriptomes of prostates from wild-type mice to those from littermates isolated 3 days post-castration (Table S1). In parallel, we compared transcriptome data from prostates isolated from intact Pten+/+ (wild-type) and Pten−/− mice (without castration). GSEA revealed that genes up- or downregulated in response to castration in wild-type mice were significantly enriched in intact Pten−/− prostates compared to intact Pten+/+ prostates, indicating that Pten loss is associated with reduced AR activity (Figure 3E). Examination of individual genes revealed that a substantial number of the genes up- or downregulated by castration in intact mice are already up- or downregulated in intact Pten−/− mice (Figure S2C and Table S2). Together with the human prostate tumor data and the BEZ235 treatment studies, these findings establish that the increase in PI3K activation associated with PTEN loss impairs AR signaling.

**Inhibition of AR Promotes PI3K Activity in PTEN-Loss Prostate Cancer**

Previous studies in mouse models and cell lines have implicated PTEN loss as a potential cause of castration resistance (Gao et al., 2006; Zhang et al., 2009). Our finding that PI3K activation is associated with reduced AR output suggests a potential explanation, e.g., these tumors are less dependent on AR. However, it is possible that AR function, albeit low, remains intact due to low-circulating androgens that remain after castration. To investigate the potential role of persistent AR signaling in this context, we evaluated the effect of combined androgen blockade (surgical castration plus the next generation AR inhibitor MDV3100 that impairs AR nuclear translocation and DNA binding) (Tran et al., 2009) in the Pten−/− model. After 7 days of treatment, mRNA levels of the androgen-regulated genes Pbsn, Nkx3.1, and Pscα were decreased 25- to 50-fold (Figure S3A), and AR protein levels were primarily cytoplasmic (Figure 4B), confirming substantial inhibition of AR pathway output in tumors isolated from treated mice. Despite this magnitude of pathway inhibition, tumors showed only modest regression (mean reduction in tumor volume of 19.2%) without obvious histologic changes (Figures 4A and 4B; Figure S3B). In addition there was minimal effect on proliferation as measured by Ki67 staining (proliferative indices of 0.34 and 0.28 pretreatment and posttreatment, respectively) (Figure 4B; Figure S3D). In contrast the same treatment regimen in PB-MYC mice resulted in profound reductions in tumor volume (mean of 81.4%), near-complete pathologic responses, and virtually absent Ki67 staining (reduction in proliferative index from 0.554 to 0.012; p < 0.001) (Figure 4A; Figures S3B–S3D). We conclude that even combined AR blockade remains ineffective in Pten−/− mice.

Although it is formally possible that the 50-fold impairment in AR output was simply not enough to impair survival of PTEN-deficient prostate cells, another explanation could be persistent survival signaling through AKT. Remarkably, AKT phosphorylation at Ser473 was increased in prostates of Ptenlox/lox mice following castration. This increase was likely PI3K pathway dependent because it was inhibited by concurrent treatment with BEZ235 (Figure 4C). Similar results, including increased phosphorylation of downstream AKT targets such as GSK-alpha and PRAS40, were observed in PTEN-negative LNCaP cells treated with MDV3100 (Figure 4D). We also observed increased levels of pAKT in the AR-positive cell line LAPC4 following treatment with MDV3100 (Figure S3E). The effects of MDV3100 on AKT activation are likely specific to AR inhibition because siRNA knockdown of AR gave similar results (Figure 4D), and no change in pAKT levels was observed in AR-negative PC3 cells (Figure S3F).

The immunophilin FKBP5 is a chaperone for the AKT phosphatase PHLPP, and its expression in prostate cancer is androgen dependent (Gao et al., 2005; Pei et al., 2009). We hypothesized that AR inhibition would result in reduced FKBP5 expression and, consequently, lower PHLPP protein levels, and this could cause increased phosphorylation of AKT. Indeed, FKBP5 and PHLPP protein levels were both reduced in LNCaP cells treated with MDV3100 or siRNA AR, and this was accompanied by an increase in phosphoAKT (Figure 4D). siRNA knockdown of PHLPP in the LNCaP cell line resulted in increased levels of pAKT as expected, and importantly, knockdown of FKBP5 resulted in decreased levels of PHLPP and upregulation of pAKT, phenocopying the effects of MDV3100 (Figure 4E; Figure S3G). Furthermore, constitutive expression of FKBP5 (CMV-FKBP5) resulted in stable levels of PHLPP and blocked the upregulation of pAKT in the presence of MDV3100 (Figure 4E). Protein levels of PHLPP were also lower in Ptenlox/lox mice following castration (Figure 4F). These data suggest that AR negatively regulates AKT activity through stabilization of PHLPP. Consequently, AR inhibition destabilizes PHLPP and results in unchecked AKT activation, especially in the setting of PTEN loss.

**Combined PI3K and AR Pathway Inhibition Gives Profound Tumor Regressions**

Taken together, the effects of PI3K inhibitors on the AR pathway and AR inhibitors on the PI3K pathway in PTEN-deficient prostate cells demonstrate that perturbations in the activity of one pathway impact signaling through the other pathway. Therefore, we evaluated the effect of combined PI3K and AR pathway inhibition in PTEN-deficient LNCaP cells and in the conditional Pten−/− prostate cancer model. BEZ235 and MDV3100 each displayed modest single-agent antiproliferative activity in LNCaP cells (primarily cytostatic), but neither treatment promoted
apoptotic cell death (Figures 5A and 5B). However, the combination of BEZ235 with MDV3100 led to a profound decrease in cell number and an increase in cleaved PARP, a marker of apoptosis (Figures 5A and 5B). To determine if similar effects might be observed by inhibiting mTORC1 or MEK, we compared the effects of RAD001 or PD0325901 to BEZ235, alone and in various combinations, including with MDV3100 (Figures S4A and S4B). The greatest antiproliferative effect was observed with combined treatment with BEZ235 and MDV3100, indicating that PI3K and/or mTORC1/2 and AR, but not mTORC1 or MEK, appear to be the most critical targets in this model. Based on our discovery that inhibition of the PI3K pathway promotes AR activity in a HER2/3-dependent manner, we reasoned that a HER2/3 inhibitor might be similarly efficacious (replacing MDV3100) in combination with BEZ235. Indeed, combined treatment with BEZ235 and PKI166 was as effective as BEZ235 plus MDV3100 (Figures 5C and 5D). Furthermore, inhibition of HER2/3 abolished the upregulation of AR protein

Figure 4. Preclinical Trial of Combined Androgen Blockade in GEM Models of Prostate Cancer

(A) Waterfall plot depicting proportional change in tumor response for PB-MYC (n = 5) and Pten\textsuperscript{lox/lox} (n = 5) mice treated with castration and MDV3100 (30 mg/kg/day).

(B) Histologic and immunohistochemical staining for AR and ki67 and Pten\textsuperscript{lox/lox} mice treated with vehicle control or castration and MDV3100 (microscopy at 200X).

(C) Western blot analysis of prostates from wild-type and Pten\textsuperscript{lox/lox} mice treated with surgical castration and/or BEZ235 for 7 days. Ventral prostates from individual 8-week-old mice were analyzed.

(D) Western blot analysis for PHLLPP, FKBP5, and pAKT in LNCaP treated with MDV3100 (10 \mu M) or siRNA AR for 24 hr.

(E) Western blot analysis for PHLLPP, FKBP5, and pAKT in LNCaP transfected with siRNA FKBP5. Western blot analysis for PHLLPP, FKBP5, and pAKT in LNCaP transfected with vector control, treated with MDV3100 (10 \mu M), or transfected with CMV-FKBP5 and treated with MDV3100.

(F) PHLLPP immunohistochemistry of intact and castrate Pten\textsuperscript{lox/lox} mice. See also Figure S3.
levels, and transcriptional activity observed with PI3K pathway inhibition (Figure S4C), as measured by PSA expression.

To test the impact of combined PI3K/AR therapy in tumor models, \(Pten^{\text{lox/lox}}\) mice with established prostate tumors were treated with BEZ235 plus MDV3100 and castration. Combined PI3K and AR pathway inhibition led to dramatic reductions in tumor volume (mean tumor volume reduction of 84.25%), with near-complete pathologic responses and no evidence of residual cell proliferation detectable by Ki67 staining (Figures 5E and 5F; Figure S4C). Combined PI3K/AR therapy also induced regressions in LNCaP xenografts (24% mean reduction in tumor volume), whereas average tumor volume in mice treated with vehicle or single-pathway therapy (BEZ235 or castration) increased (28%, 30%, and 31%, respectively) (Figure S4D). Addition of BEZ235 to castration plus MDV3100 in PB-MYC mice showed no measurable benefit, but the substantial response to combined androgen blockade alone in this model makes it difficult to detect any effect of combined PI3K/AR therapy (Figure 5F).
DISCUSSION

AR pathway inhibition has long been the treatment of choice for men with metastatic prostate cancer. Although much attention has been devoted to mechanisms of acquired resistance, there has been little investigation of the considerable variability in primary response among patients. Here, we show, by mRNA transcriptome analyses, that activation of the PI3K pathway is associated with repressed androgen signaling in mouse and human prostate cancers and that this may, in part, be responsible for the castrate-resistant phenotype observed with these prostate tumors. Importantly, we demonstrate that this resistance is reversible because inhibition of the PI3K pathway restores AR signaling in PTEN-deficient prostate cells. At least one mechanism appears to be through relief of negative feedback to HER kinases. Similarly, blockade of AR relieves feedback inhibition of AKT by the phosphatase PHLPP. This reciprocal feedback regulation of the PI3K and AR pathways provides a compelling explanation for the poor efficacy of single-pathway therapy in PTEN null cancers and the substantially better effects of combined PI3K/AR pathway inhibition (Figure 5G).

Prior work has implicated PTEN loss as a potential cause of castration resistance in mice and in humans (Gao et al., 2006; Zhang et al., 2009). Zhang et al. (2009) reported that Pt en prostate conditional null mice treated with surgical castration have a delay in tumor growth and minimal tumor regression. Although, to our knowledge, no human studies have formally addressed this question, there is evidence from presurgical treatment studies that tumors with PTEN loss are relatively refractory to bicalutamide (Ham et al., 2009; Schulman et al., 2000). Despite the evidence that PTEN loss can promote castration resistance, there is little insight into the mechanism. Some reports have suggested that PTEN loss activates AR, through PI3K-mediated stabilization of AR protein levels or AKT-mediated phosphorylation and transcriptional activation of AR. Conversely, other studies have demonstrated that PI3K activation promotes degradation of AR and inhibits AR transcriptional activity (Lin et al., 2003, 2004; Wang et al., 2008). Our transcriptome studies make a strong case for the latter model. In addition our finding that reduced expression of the AR target gene FKBP5 results in an increase in AKT activation (via PHLPP degradation) in PTEN null cancers further explains the survival advantage of these tumor cells in the setting of castration.

This work has immediate implications for the design of clinical trials evaluating PI3K pathway inhibitors in prostate cancer. Our preclinical data predict that single-agent PI3K pathway inhibitor therapy will most likely result in disease stabilization rather than tumor regression, particularly in PTEN null tumors that represent ~40% of primary cancers and ~70% of metastases (Taylor et al., 2010). Additionally, given that the primary serum marker used to monitor disease progression (prostate-specific antigen or PSA) is androgen regulated, patients treated with PI3K pathway inhibitors may experience a rise in PSA level if their tumors are PTEN deficient. Our data argue that combined therapy with an AR pathway inhibitor is required for maximal efficacy in PTEN null cancers. In patients with hormone-naive disease, this could be achieved using currently available antiandrogen therapy, but patients with castration-resistant prostate cancer are likely to require next-generation AR pathway inhibitors such as abiraterone or MDV3100.

Because BEZ235 inhibits both PI3K and mTORC1/2, our data do not delineate which target is most critical for the observed effects of combination therapy. Others reported beneficial effects of combined AR and mTORC1 (rapalogs) inhibition in a similar Ptenlox/lox model, but the magnitude of tumor response was less substantial because mice had significant amounts of residual tumor tissue at the time of sacrifice (Zhang et al., 2009). In addition these investigators monitored tumor volume by ultrasound, which makes it difficult to distinguish between shrinkage caused by true tumor regression versus a reduction in the cystic dilation that accompanies Pten+/− prostate tumors. Kinkade et al. (2008) also reported benefit from combining rapamycin with a MEK inhibitor in Nkx3.1−/−;Pten+/− mice, but this experiment differs in that Pten+/− mice have a less aggressive cancer phenotype than the Ptenlox/lox model. Side-by-side experiments using identical endpoints in the same model are required to properly compare these regimens. In the meantime our in vitro studies establish that dual PI3K/mTORC1/2 inhibition (BEZ235) is superior to mTORC1 inhibition (RAD001) when combined with AR blockade and that MEK inhibition (PD0325901) is relatively ineffective. Because BEZ235 inhibits mTORC1/2 more potently than PI3K, it is possible that the superiority of BEZ235 over RAD001 is solely through TORC1/2 blockade (Serra et al., 2008). This question can be addressed using selective TORC1/2 inhibitors (Chresta et al., 2010; Thoreen et al., 2009).

Our finding that HER2/3 activation (and subsequent AR activation) is associated with PI3K pathway inhibition also has important clinical implications because a HER2 kinase inhibitor such as lapatinib could, in theory, replace the requirement for an antiandrogen in combination with a PI3K pathway inhibitor. Our studies with the preclinical HER2 inhibitor PK1-166 establish this principle in vitro. Single-agent trials with HER2 inhibitors (trastuzumab, pertuzumab, lapatinib) in men with castration-resistant prostate cancer have been largely negative (Solit and Rosen, 2007), but our data suggest that combination of these inhibitors with PI3K pathway inhibitors is required to elicit activity.

In summary our results demonstrate that inhibition of the PI3K pathway in Pten−/− prostate cancer results in feedback signaling to the receptor tyrosine kinase HER2/HER3 leading to activation of AR. Conversely, blockade of AR results in activation of AKT through reduced levels of FKBP5 impairing the stability of PHLPP. This bidirectional crosstalk between two critical survival pathways in prostate cancer provides the molecular rationale for simultaneously targeting both pathways. The success of clinical trials evaluating PI3K pathway inhibitors in prostate cancer could be optimized by enrolling patients with documented activation of the PI3K pathway and treating in combination with appropriate AR pathway inhibition.

EXPERIMENTAL PROCEDURES

Mouse Models of Prostate Cancer

Animal studies were carried out under protocol 06-07-012 approved by the MSKCC Institutional Animal Care and Use Committee. Institutional guidelines for the proper, humane use of animals in research were followed. The GEM

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models of human prostate cancer (PB-MYC and Pten\textsuperscript{Cre/lox} PB-Cre) have been described previously (Elwood-Yen et al., 2003; Trotman et al., 2003). Genotyping was conducted through our core facility using previously published primer sets and protocols. PB-MYC (ages 12 and 18 months) and Pten\textsuperscript{lox} (age 6 months) were imaged by our MRI small animal imaging core prior to and at the completion of treatment (see Supplemental Experimental Procedures) (Chen et al., 2005). Surgical castration was performed under anesthesia with isoflurane. Mice were monitored postoperatively for recovery from anesthesia and checked daily for 2 days postoperatively. Surgical skin clips were removed on postoperative day 5. Mice undergoing treatment were administered control vehicle or therapeutic doses of the appropriate agents by oral gavage on a Monday through Friday schedule for a total of 35 days. Any mouse suffering distress or greater than 15% weight loss during treatment was euthanized by CO\textsubscript{2} asphyxiation. MRI tumor volumes were reported for each mouse at time point zero (T0) at initiation of study and time point 35 days (T35) at completion of study. Changes in tumor volumes between T0 and T35 were calculated for individual mice and reported in waterfall plots. At the completion of study, mice were euthanized by CO\textsubscript{2} asphyxiation, and tissue was procured for histology, mRNA analysis, protein analysis, and tissue banking.

For xenograft experiments, \(1 \times 10^7\) LNCaP cells were injected into the bilateral flanks of SCID mice. When mice tumors were approximately 500 mm\(^3\), mice were randomized to the treatment groups. Tumor volume was measured bimonthly for a total of 2 weeks, and the animals were sacrificed according to our protocol. All animal experiments conform to the relevant regulatory standards and were approved by our IACUC committee under our approved animal protocol.

**Targeted Pathway Inhibitors**

The AR inhibitor MDV3100 was synthesized by the MSKCC chemistry core and used in vitro at a concentration of 1 \(\mu\)M and in vivo with a dose of 30 mg/kg/day administered once daily by oral gavage on a Monday through Friday schedule. The PI3K pathway inhibitors NVP-BEZ235 (PI3K and mTORC1/2 inhibitor) and RAD001 (mTORC1 inhibitor) were provided by Novartis under a Materials Transfer Agreement. The concentration of BEZ235 and RAD001 used for in vitro experiments was 500 and 100 \(\mu\)M, respectively. For in vivo experiments the dose of BEZ235 used was 45 mg/kg/day administered once daily by oral gavage on a Monday through Friday schedule. The HER2 kinase inhibitor PK166 was provided by Novartis and used for in vitro experiments at a concentration of 5 \(\mu\)M. PD0325901 (MEK inhibitor) was synthesized by the MSKCC chemistry core and used for in vitro studies at a concentration of 1 \(\mu\)M. AKT1/2 inhibitor (AKT VIII) was purchased from Calbiochem and used in vitro at a concentration of 1 \(\mu\)M.

**Mouse mRNA Expression Analysis**

Prostate tissues frozen for total RNA isolation were homogenized in TRIzol Reagent (Invitrogen; #15596-026), followed by phase separation, washing, precipitation, and resuspension of RNA in RNase-free water according to manufacturer’s protocols. The RNA was further purified using the RNeasy kit (QIAGEN) according to manufacturer's protocols, followed by quantification and normalization using A\textsubscript{260}/A\textsubscript{280}. cDNA synthesis from 1 \(\mu\)l of cDNA in a total volume of 20 \(\mu\)l was carried out using the TaqMan Reverse Transcription Reagents (Applied Biosystems) with random hexamers according to the manufacturer’s protocol. Triplicate samples for quantitative PCR were run in the realaplex Mastercycler (Eppendorf) using the Power SYBR Green PCR Mastermix (Applied Biosystems). Each reaction contained 1 \(\mu\)l of cDNA in a total volume of 20 \(\mu\)l. \(\lambda\Delta C\text{t}\) for each gene was determined after normalization to Hprt, and \(\Delta C\text{t}\) was calculated relative to the designated reference sample. Gene expression values were then expressed as a fold change, calculated by \(2^{-\Delta\Delta C\text{t}}\). See Supplemental Experimental Procedures for primer sequences.

**Mouse Microarray Expression Profiling**

Microarray gene expression profiling was performed on RNA prepared from the prostates of wild-type and Pten\textsuperscript{Cre/lox} PB-Cre intact and castrate mice. Eight-week-old wild-type (\(n = 7\)) and Pten prostate conditional null (\(n = 7\)) mice in the C57B6 background were used. Three mice of each genotype were castrated. Three days after castration, mice were euthanized, and RNA was isolated from prostates then profiled on the Illumina MouseRef-8 v2 bead arrays. Raw data were imported into Partek Genomics Suite v6.5 where data were Log2 transformed and quartile normalized. See Supplemental Experimental Procedures for detailed methods for generation of murine androgen-responsive gene signature and GSEA.

**In Vitro Experiments**

In vitro experiments were conducted using the LNCaP and PC3 cell lines obtained from American Type Culture Collection. LAPC4 and LNCaP AR-ARE-Luciferase, which expresses exogenous AR and Luciferase expression under control of an androgen-regulated promoter, have been previously described (Tran et al., 2009). Proliferation assays were conducted by plating \(1 \times 10^5\) cells per well of a 12-well cell culture plate and treating with vehicle (DMSO) control or AR/PI3K inhibitors at the aforementioned concentrations. Viable cells were counted using a hemocytometer using trypan blue exclusion on days 1, 3, and 5. Cell lysates for western blot analysis were prepared using standard RIPA buffer. Luciferase assays were conducted using the Promega ONE-Glo Luciferase Assay System and measured using a luminometer plate reader. All in vitro experiments were conducted in triplicate, and standard deviations were reported. Significance was determined by the Student’s t-test. The FKB5P, PHLPP, AKT1, AKT2, and AR siRNA SMARTpool was obtained from Dharmacon. Control siRNA luciferase was used for all experiments. The CMV-FKB5P plasmid was purchased from OriGene.

**Immunohistochemical and Western Blotting Antibodies**

The antibodies used for western blot analysis and immunohistochemistry were pAKT Ser473 (Cell Signaling Technology; 1:1000 dilution), pAKT Thr308 (Cell Signaling Technology; 1:1000 dilution), AKT (Cell Signaling Technology; 1:1000 dilution), pERK Thr202/Tyr204 (Cell Signaling Technology; 1:1000 dilution), ERK (Cell Signaling Technology; 1:1000 dilution), pPRAS40 Thr246 (Cell Signaling Technology; 1:1000 dilution), PRAS40 (Cell Signaling Technology; 1:1000 dilution), pGSK3-a Ser21 (Cell Signaling Technology; 1:1000 dilution), GSK (Cell Signaling Technology; 1:1000 dilution), PARP (Cell Signaling Technology; 1:1000 dilution), AR N-20 (Santa Cruz Biotechnology; 1:500 dilution), c-MYC (Epitomics; 1:500 dilution), PHLPP (Bethyl Laboratories; 1:500 dilution), FKB5P (Bethyl Laboratories; 1:500 dilution), HER2 (Cell Signaling Technology; 1:1000 dilution), HER3 (Cell Signaling Technology; 1:1000 dilution), HER2 (Cell Signaling Technology; 1:1000 dilution), and Actin (Cell Signaling Technology; 1:1000 dilution). All immunohistochemical analyses were conducted by the MSKCC Molecular Cytology Core.

**Human CGH and mRNA Profiling**

Our human prostate cancer data set has been previously published (Taylor et al., 2010). All patients provided informed consent, samples were procured, and the study was conducted under MSKCC Institutional Review Board approval. Briefly, copy number data were generated on Agilent 244K aCGH arrays, and mRNA expression data were obtained on Affymetrix Human Exon 1.0 ST arrays. The complete genomics data set and analytic methods are reported separately and are available at: http://cibio.mskcc.org/cancergenomics-dataportal/.

**Pathway Analysis**

The raw and normalized microarray data have been deposited into the NIH NCBI Gene Expression Omnibus (GEO), GSE24691: http://www.ncbi.nlm.nih.gov/geo/query/acc.cgi?token=keffccomgsvs&acc=GSE24691.

**ACCESSION NUMBERS**

The raw and normalized microarray data have been deposited into the NIH NCBI Gene Expression Omnibus (GEO), GSE24691: http://www.ncbi.nlm.nih.gov/geo/query/acc.cgi?token=keffccomgsvs&acc=GSE24691.
SUPPLEMENTAL INFORMATION

Supplemental Information includes four figures, Supplemental Experimental Procedures, and two tables and can be found with this article online at doi:10.1016/j.ccr.2011.04.008.

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Death from prostate cancer. Br. J. Cancer


