

The PI3K Pathway As Drug Target in Human Cancer

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ABSTRACT

The phosphatidylinositol 3-kinase (PI3K) signaling axis impacts on cancer cell growth, survival, motility, and metabolism. This pathway is activated by several different mechanisms in cancers, including somatic mutation and amplification of genes encoding key components. In addition, PI3K signaling may serve integral functions for noncancerous cells in the tumor microenvironment. Consequently, therapeutics targeting the PI3K pathway are being developed at a rapid pace, and preclinical and early clinical studies are beginning to suggest specific strategies to effectively use them. However, the central role of PI3K signaling in a large array of diverse biologic processes raises concerns about its use in therapeutics and increases the need to develop sophisticated strategies for its use. In this review, we will discuss how PI3K signaling affects the growth and survival of tumor cells. From this vantage, we will consider how inhibitors of the PI3K signaling cascade, either alone or in combination with other therapeutics, can most effectively be used for the treatment of cancer.

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INTRODUCTION

It has been more than 20 years since phosphatidylinositol 3-kinase (PI3K) was first discovered. The transforming ability of viral oncoproteins relied on an association with a PI3K lipid kinase activity.¹⁻⁴ Over the ensuing years, studies established the central role of PI3K signaling in several cellular processes critical for cancer progression, including metabolism, growth, survival, and motility. Inappropriate co-option of PI3K signaling is one of the most frequent occurrences in human cancer.^{5,6} Consequently, significant efforts have been made to generate inhibitors of the PI3K pathway to treat cancers. However, it remains unknown which cancers will benefit most from these treatments and how to best use such therapeutics. In addition, the many possible untoward biologic sequelae of PI3K inhibition may limit the potential therapeutic gain of PI3K pathway inhibition. Here we will review data demonstrating the role of PI3K in tumor development and maintenance. We will compare the different potential therapeutic options for inhibiting this pathway and how their efficacy may be affected by the mechanism of PI3K pathway activation in a particular cancer. Finally, we will discuss the emerging data assessing the relative benefits of PI3K pathway inhibitors used as single agents versus combination therapies to treat cancer.

PI3K SIGNALING CASCADE REGULATES CELL GROWTH AND SURVIVAL

There are three classes of PI3Ks grouped according to structure and function. Class I_A PI3K is the one most clearly implicated in human cancer.⁷ Class I_A PI3Ks consist of a regulatory subunit and a catalytic subunit. Three mammalian genes, *PIK3R1*, *PIK3R2*, and *PIK3R3*, encode p85 α (p85 α , p55 α , and p50 α isoforms), p85 β , and p55 γ regulatory subunits, respectively, which by convention are referred to collectively as p85.^{5,7,8} The catalytic isoforms, p110 α , p110 β , and p110 δ , are the products of three genes, *PIK3CA*, *PIK3CB*, and *PIK3CD*.^{5,8} As will be discussed in greater detail below, both *PIK3CA* and *PIK3R1* are somatically mutated in cancers, and these mutations promote activation of the PI3K pathway.⁹⁻¹²

Class I_A PI3Ks are activated by growth factor stimulation through receptor tyrosine kinases (RTKs).¹³⁻¹⁵ The regulatory subunit, p85, directly binds to phosphotyrosine residues on RTKs and/or adaptors.¹⁶ This binding relieves the intermolecular inhibition of the p110 catalytic subunit by p85 and localizes PI3K to the plasma membrane where its substrate, phosphatidylinositol 4,5-bisphosphate (PI[4,5]P₂), resides.^{15,16} PI3K can also be stimulated by activated Ras, which directly binds p110.¹⁷ Additionally, the p110 β catalytic subunit can be activated by G-protein coupled receptors.⁸

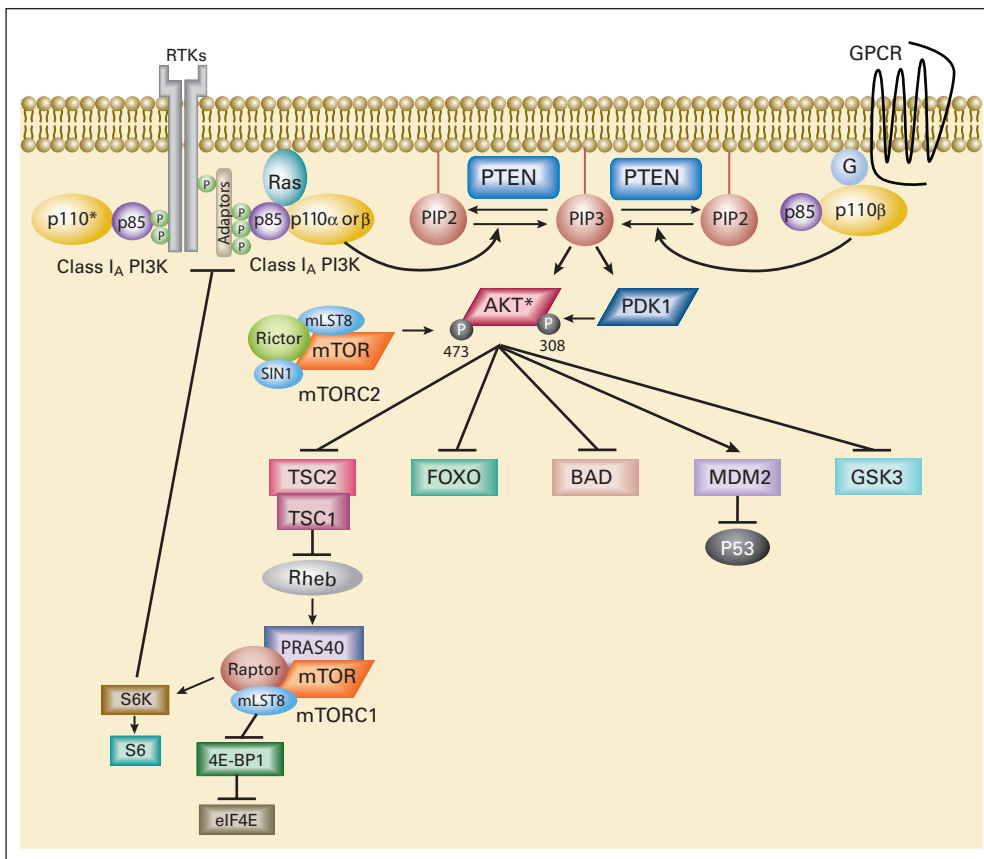


Fig 1. The phosphatidylinositol 3-kinase (PI3K) signaling cascade. PI3K signaling impacts on cell growth, survival, and metabolism. Arrows represent activation, while bars reflect inhibition. A negative feedback loop has been described from the downstream target S6 kinase (S6K) to the adaptor protein IRS-1. RTK, receptor tyrosine kinase; GPCR, G-protein coupled receptor; P, phosphate; G, G protein; PTEN, phosphatase and tensin homolog; IRS-1, insulin receptor substrate 1; eIF4E, eukaryotic initiation factor 4E; S6, ribosomal S6 protein; PIP₂, phosphatidylinositol 4,5-bisphosphate; mTORC2, rapamycin (mTOR) –containing protein complex 2. (*) p110 alpha, beta, or delta.

PI3K phosphorylates PIP₂ on the 3'OH position to produce PI(3,4,5)P₃ (PIP₃; Fig 1). The tumor suppressor phosphatase and tensin homolog deleted on chromosome 10 (PTEN) dephosphorylates PIP₃ to PIP₂, thereby terminating PI3K-dependent signaling. PIP₃ propagates intracellular signaling by directly binding pleckstrin homology (PH) domains of various signaling proteins.¹⁸ PI(3,4,5)P₃ brings two PH domain-containing serine/threonine kinases, phosphoinositide-dependent kinase 1 (PDK1) and AKT, into close proximity. PDK1 activates AKT by phosphorylating AKT at threonine 308.^{19–21} PI3K-AKT signaling promotes cell growth and survival by several mechanisms. AKT promotes cell survival by inhibiting proapoptotic Bcl-2 family members BAD and BAX.^{5,18} AKT also impedes negative regulation of the transcription factor NF- κ B, leading to increased transcription of antiapoptotic and prosurvival genes.²² Phosphorylation of Mdm2 by AKT antagonizes p53-mediated apoptosis, and AKT negatively regulates forkhead transcription factors, thereby reducing production of cell death-promoting proteins.²² AKT also phosphorylates TSC2, thereby inhibiting the rheb GTPase activity of the TSC1/TSC2 dimer. Activated rheb stimulates the mammalian target of rapamycin (mTOR) –containing protein complex mTORC1, leading to increased p70 S6 kinase activity.⁵ Activation of mTORC1 results in increased protein synthesis by phosphorylation of eukaryotic initiation factor 4E and the ribosomal S6 protein.⁵ While mTORC1 relays signals following PI3K-AKT activation, a second mTOR complex, mTORC2, contributes to complete AKT activation by phosphorylating AKT on serine 473.^{23–25} Of note, activation of the mTORC1 target, S6 kinase, negatively feeds back to diminish PI3K

activation. S6 kinase can phosphorylate and inhibit the adaptor protein insulin receptor substrate 1, thereby inhibiting insulin or insulin-like growth factor 1–mediated PI3K activation.^{26–28}

Inhibitors of PI3K Signaling in Cancer Treatment

Inhibition of PI3K signaling can diminish cell proliferation, and in some circumstances, promote cell death. Consequently, components of this pathway present attractive targets for cancer therapeutics. A number of PI3K pathway inhibitors have been developed and are being evaluated in preclinical studies and in early clinical trials. Rapamycin analogs, such as temsirolimus and everolimus, that specifically inhibit mTORC1 are the most advanced in the clinic, and they have received US Food and Drug Administration approval for the treatment of advanced renal cell carcinoma.²⁹ The role for rapamycin analogs in the treatment of cancer has been extensively reviewed elsewhere and thus will not be discussed further.³⁰ In this review, we will discuss the potential therapeutic roles for other PI3K pathway inhibitors. These include PI3K inhibitors (both pan-PI3K and isoform-specific PI3K inhibitors), dual PI3K-mTOR inhibitors that are catalytic site inhibitors of the p110 isoforms and mTOR (the kinase component of both mTORC1 and mTORC2), mTOR catalytic site inhibitors, and AKT inhibitors. Not only do these agents have the capacity to inhibit cancer cell proliferation and survival signals as described above, but they may also impact tumor angiogenesis, metastasis, and metabolism. Due to space limitations, the impact of PI3K inhibition on tumor angiogenesis and cell motility is discussed in the Appendix (online only).

PI3K and Insulin Signaling: Potential Toxicity and Pharmacodynamic Marker of PI3K Inhibition

PI3K signaling has a central role in mediating the effects of insulin on cellular metabolism that is conserved throughout eukaryotic evolution.⁵ Noninsulin-dependent diabetes mellitus, marked by insulin insensitivity, is associated with diminished PI3K response to insulin signaling.^{5,31} Several transgenic and knockout mice harboring alterations in p85, p110, PTEN, or AKT2 validate the functional significance of this pathway on glucose homeostasis.³¹⁻³⁴ These data suggest that insulin resistance may be observed in patients treated with PI3K pathway inhibitors, and indeed this may be used as a pharmacodynamic marker of target inhibition in patients. As will be discussed further below, initial phase I studies with PI3K pathway inhibitors have demonstrated some signs of insulin resistance, but this has not been a dose-limiting toxicity. While both p110 α and p110 β appear to play specific roles in insulin signaling, studies suggest that glucose homeostasis is predominantly mediated by p110 α .^{35,36} Inhibitors of p110 α , but not p110 β or p110 δ , have been shown to inhibit insulin-stimulated glucose uptake in adipocytes and to block insulin-mediated glucose regulation in mice.³⁶ Consequently, in settings where p110 β appears to be the critical PI3K catalytic isoform mediating transformation (eg, some PTEN-deficient tumors, see below), a p110 β -specific inhibitor may offer efficacy with decreased risk of insulin resistance compared with a pan-PI3K inhibitor.

ACTIVATION OF PI3K SIGNALING IN CANCER

PI3K signaling is activated in human cancers via several different mechanisms.^{6,11-13} Increased PI3K signaling is often due to direct mutational activation or amplification of genes encoding key components of the PI3K pathway such as *PIK3CA* and *AKT1*, or loss of *PTEN*.^{6,9,12,37-41} Genetic alterations in several components of the PI3K

signaling pathway have been reported and are summarized in Table 1. PI3K also can be activated by genetic mutation and/or amplification of upstream RTKs, and possibly by mutationally activated Ras.^{7,17} The mechanism of PI3K activation in an individual cancer may suggest the most effective type of therapeutic to inhibit the pathway.

Somatic Alterations of PI3K Pathway Components in Cancer

The most common genetic alteration of the PI3K signaling pathway found in human cancer is inactivation of the *PTEN* tumor suppressor gene. Inactivation of *PTEN* leads to loss of its lipid phosphatase activity, causing accumulation of PIP₃.^{56,57} The majority of somatic mutations in *PTEN* lead to protein truncation. However, missense mutations that typically abrogate PIP₃ phosphatase activity are also commonly noted.⁵⁸ While most *PTEN* mutations are sporadic, germline mutations in *PTEN* are noted in hereditary neoplastic disorders, such as Cowden disease.⁵⁹ Homozygous and hemizygous deletions of *PTEN* are also observed in human cancers.^{38,45} Transcriptional repression and epigenetic silencing of *PTEN*, typically through promoter hypermethylation, is also a mechanism of *PTEN* inactivation.^{42,43} Because there are both genetic and epigenetic causes for *PTEN* loss, accurate assessment of a cancer's *PTEN* status remains challenging and may ultimately require reliable measurements of protein expression.

More recently, somatic mutations in *PIK3CA* have been identified in a variety of human tumors, including breast, colon, and endometrial cancers and glioblastomas (see Catalogue of Somatic Mutations in Cancer, <http://www.sanger.ac.uk/genetics/CGP/cosmic>).^{6,7,9,12,39} Most of these mutations cluster to two hot spot regions in exons 9 and 20.^{6,12} Exon 20 encodes the catalytic domain of p110 α , and mutations in this domain may constitutively activate its enzymatic activity. Exon 9 encodes the helical domain of p110 α , and

Table 1. Genetic Alterations of the PI3K Signaling Pathway in Cancer

Gene	Alteration	Comments	Tumor Types	References
<i>PTEN</i>	LOF mutation	Truncation; loss of phosphatase activity	Bladder, brain, breast, cervical, colorectal, endometrial, gastric, head and neck, kidney	COSMIC, Li et al ³⁸
	Deletion	Homozygous or hemizygous	Leukemia, liver, lung, lymphoma, melanoma, ovary, prostate, thyroid	
	Epigenetic silencing	Transcriptional repression, usually by promoter hypermethylation	Breast, colon, melanoma	Garcia et al, ⁴² Goel et al, ⁴³ Berns et al ⁴⁴
<i>PIK3CA</i>	GOF mutation	Exon 9 (E542K, E545K) helical domain; exon 20 (H1047R) catalytic domain	Breast, colorectal, glioblastoma, endometrial, cervical, esophageal, gastric, head and neck, liver, lung, lymphoma, ovarian, pancreatic, prostate, thyroid	COSMIC, Samuels et al, ⁶ Samuels et al ¹²
	Amplification	Increased protein levels and activity	Breast, cervical, gastric, lung, ovarian, prostate	Sun et al, ⁴⁵ Byun et al, ⁴⁶ Campbell et al, ⁴⁷ Ma et al, ⁴⁸ Rácz et al ⁴⁹
<i>PIK3R1</i>	GOF mutation	Loss of C-terminal inhibitory domain; constitutive activity	Brain, colon, ovarian	Mizoguchi et al, ¹⁰ Philp et al ¹¹
<i>AKT1</i>	GOF mutation	Pleckstrin homology domain, membrane localization, and constitutive activation	Breast, colon, endometrial, melanoma, ovarian	Shoji et al, ⁴⁰ Carpten et al, ⁵⁰ Davies et al ⁵¹
<i>AKT2</i>	GOF mutation	Kinase domain mutation	Colorectal	Parsons et al ⁵²
	Amplification		Breast, colon, lymphoma, pancreas	Cheng et al, ⁵³ Cheng et al, ⁵⁴ Bellacosa et al ⁵⁵
<i>AKT3</i>	GOF mutation	Pleckstrin homology domain, membrane localization, and constitutive activation	Melanoma	Davies et al ⁵¹
<i>PDK1</i>	GOF mutation	Kinase domain mutation	Colorectal	Parsons et al ⁵²

Abbreviations: LOF, loss of function; COSMIC, Catalogue of Somatic Mutations in Cancer at www.sanger.ac.uk/genetics/CGP/cosmic; GOF, gain of function.

these mutations de-repress an inhibitory interaction between the N-terminal SH2 domain of p85 and the p110 α catalytic subunit.^{60,61} A smaller cluster of mutations is also found in the N-terminal p85 interacting domain. Interestingly, these mutations increase the lipid kinase activity of p110 α but do not appear to alter the interaction between p110 α and p85 α .^{9,12}

Expression of these *PIK3CA* mutants leads to increased oncogenic potential in vitro and in vivo.^{9,37} They cause constitutive signaling along the PI3K pathway in the absence of growth factors and therefore seem to obviate the usual obligate interactions with tyrosine phosphorylated RTKs and/or adapters. Thus, it is intriguing that some studies have suggested that the presence of these mutations confers resistance to therapies targeting RTKs.^{44,62} Expressing mutated *PIK3CA* in fibroblasts and mammary epithelial cells results in transformation, growth factor-independent proliferation, and resistance to apoptosis.^{9,63,64} Additionally, transgenic mice with lung-specific induction of the kinase-domain mutant p110 α H1047R develop lung adenocarcinomas.⁶⁵ In addition to these activating mutations, amplification of *PIK3CA* is also observed frequently in ovarian cancer and other tumors, but how amplification affects PI3K activation is less clear.³⁹

Mutations in the p85 regulatory subunit *PIK3R1* are also observed in a variety of human cancers, including glioblastomas, ovarian cancers, and colorectal cancers.^{10,11} Mutations in *PIK3R1* generally produce either truncations or in-frame deletions that often localize to the inter-SH2 domain of p85 α . Structural analyses suggest that the iSH2 domain of p85 interacts with the C2 domain of p110.⁶⁰ Thus, it seems likely that these p85 α mutations also activate PI3K signaling by relieving the inhibitory effect of p85 on p110.^{11,66} Laboratory studies suggest that these mutations also lead to constitutive PI3K signaling.^{11,66}

Mutations in AKT family genes encoding for AKT1, AKT2, and AKT3 have also been identified in human cancers. A single amino acid substitution, E17K, in the lipid-binding PH domain of AKT1 has been identified in breast, colorectal, endometrial, and ovarian cancers, as well as melanoma.^{40,50,51} This amino acid change alters AKT1 lipid binding, presumably leading to constitutive membrane localization in the absence of PIP₃. However, although phosphorylation on Ser473 was constitutive in this mutant, T308 phosphorylation was still responsive to PI3K activation.⁵⁰ Thus, it is unclear if PI3K inhibitors will effectively decrease AKT signaling in cancers with these mutations. The E17K mutation has also been identified in AKT3 in some melanomas.⁵¹ In addition, mutations affecting the kinase domain of AKT2 have been found in colorectal cancers; however, the functional consequences of these mutations have not been assessed.⁵² Amplification of AKT2 has also been reported in human tumors.^{53,54}

PI3K Activation by Receptor Tyrosine Kinases and Ras

In normal epithelial cells, PI3K is often activated downstream of RTK signaling. In cancers, these RTKs are often mutated, amplified, or overexpressed, causing aberrant PI3K activation. When therapies targeting RTKs are effective, they invariably lead to loss of PI3K signaling.⁶⁷ For example, PI3K is activated by epithelial growth factor receptor (EGFR) in lung cancers harboring somatic activating mutations in EGFR, and by human epidermal growth factor receptor 2 (HER2) in breast cancers with HER2 amplification.^{65,68-70} In these cancers, EGFR or HER2 phosphorylates the kinase-dead ErbB3 that, in turn, directly binds and activates PI3K.^{65,68-70} Thus, when these

cancers are successfully treated with EGFR- and HER2-targeted therapies, respectively, PI3K signaling is turned off and the cells undergo cell death. Similarly, glioblastomas frequently exhibit PI3K activation, either through integration of signaling from multiple activated RTKs, such as the constitutively active EGFRvIII mutant, or through the combined activation of RTKs and loss of PTEN.^{71,72}

The small GTPase Ras is also frequently mutated in human cancers, and PI3K is an effector of Ras-mediated oncogenic signaling.⁷³ Early studies showed that Ras directly bound p110, and provided a direct link between Ras and PI3K.⁷⁴ In addition, functional studies demonstrated that PI3K activation appears to be crucial for tumor initiation. For example, expression of a dominant-negative p85 α lacking the p110-binding domain inhibited Ras-mediated transformation.^{8,75} In addition, expression of a p110 α mutant that does not directly bind Ras inhibited K-Ras-induced lung adenocarcinomas in genetically engineered mouse models.⁷⁶ Similarly, deletion of *Pik3r1* and *Pik3r2* abrogated K-Ras G12D-induced lung tumorigenesis.⁶⁵ Although PI3K activation may be necessary for K-Ras-induced tumorigenesis, preliminary studies suggest that inhibition of PI3K signaling alone may not be sufficient to shrink established tumors in vivo or effectively treat K-Ras-mutated cancer cell lines in vitro.^{65,77} These findings underscore the difference between killing established cancers and blocking tumorigenesis and cell transformation. Furthermore, these studies suggest that established cancers with KRAS mutations may not be sensitive to single-agent PI3K pathway inhibitors.

Potential Roles for p110 β , p110 δ , and p110 γ in Transformation

While activating mutations in *PIK3CA* are frequently identified in human cancers, no oncogenic mutations have been verified in p110 β , p110 δ , or the class I_B catalytic isoform p110 γ . Although rare somatic single-residue substitutions have been found in p110 β and p110 γ (www.sanger.ac.uk/perl/genetics/CGP/cosmic), the function of these substitutions is unknown. Despite the lack of evidence for activating mutations in these other p110 catalytic isoforms, recent work has demonstrated the oncogenic potential of p110 β , p110 δ , and p110 γ . Interestingly, unlike p110 α , overexpression of wild-type p110 β , p110 δ , or p110 γ is transforming in cell culture.¹³ Although expression of the γ and δ isoforms is normally restricted to leukocytes, increased p110 δ (as well as p110 β) has been identified in some colon and bladder cancers, as well as in glioblastomas.¹³ p110 δ appears to provide the critical PI3K activity in acute myelogenous leukemia, while p110 γ is upregulated by the *Bcr-Abl* oncogene in chronic myelogenous leukemia.^{78,79}

Recent data also suggest a prominent role for p110 β in PTEN-deficient tumors. Targeted deletion of *pten* in the mouse prostate results in prostatic intraepithelial neoplasia and carcinoma.⁸⁰ Concomitant ablation of p110 β , but not p110 α , decreased PI3K proliferation signaling and prevented prostate tumorigenesis.³⁵ Similarly, inducible depletion of p110 β , but not p110 α , using short hairpin RNA in PTEN-deficient human cancer cell lines extinguished PI3K-mediated signaling and inhibited growth in vitro and in vivo.⁸¹ Deletion of p110 β also abrogated transformation of mouse embryo fibroblasts by activated Ras or EGFR mutants to a more pronounced extent than did p110 α loss.³⁵ These studies suggest that although cancers driven by *PIK3CA* mutations are candidates for treatment with p110 α -specific inhibitors, treatment of PTEN-deficient cancers may require agents with activity against p110 β .

PI3K PATHWAY INHIBITORS ENTERING THE CLINIC: PRECLINICAL AND EARLY CLINICAL DATA

A number of potential therapeutics targeting the PI3K signaling cascade have been generated. We will consider four different classes of PI3K pathway inhibitors: dual PI3K-mTOR inhibitors, PI3K inhibitors (that do not inhibit mTOR), AKT inhibitors, and mTOR catalytic site inhibitors. Table 2 summarizes PI3K pathway inhibitors in clinical trials.

Dual PI3K-mTOR Inhibitors

The catalytic domains of the p110 subunits and mTOR are structurally similar, because they all belong to the phosphatidylinositol kinase-related kinase family of kinases. Many chemical inhibitors under development inhibit both mTOR and the p110 catalytic subunits. These are termed dual PI3K-mTOR inhibitors. When compared with the other types of PI3K pathway inhibitors, dual PI3K-mTOR inhibitors have the possible advantage of inhibiting all PI3K catalytic isoforms, mTORC1, and mTORC2. Thus, they should effectively turn off this pathway completely and overcome feedback inhibition normally observed with mTORC1 inhibitors (ie, rapamycin

analogs) that may limit their efficacy (Fig 1).²⁸ However, it remains unknown if dual PI3K-mTOR inhibitors will be tolerable at doses that effectively inhibit all p110 isoforms and mTOR, or if their use will necessitate sacrificing complete inhibition of one or more of the potential targets.

For many years, the PI3K inhibitor LY294002, a dual PI3K-mTOR inhibitor, has been extensively used in preclinical studies. Although LY294002 is unsuitable for patient use, the backbone structure of this compound has been exploited in the design of novel PI3K inhibitors.^{7,90} SF-1126 (Semafore, Indianapolis, IN) is a prodrug of LY294002 that is conjugated to a tetra-peptide designed to target tumor vasculature, and this compound has demonstrated efficacy in solid tumor xenograft models.^{7,90,91} In a phase I study of SF-1126, mTORC1 inhibition in cancers was demonstrated by decreased S6 phosphorylation.⁸² No responses were observed, but stable disease was achieved in 11 of 28 patients below the maximum tolerated dose, without consistent effects on blood glucose.⁸²

Other dual PI3K-mTOR inhibitors, such as NVP-BEZ235 and NVP-BGT226 (Novartis, Basel, Switzerland) and XL765 (Exelixis, San Francisco, CA) have entered phase I testing in clinical trials.^{30,90,92} There have been several preclinical evaluations of NVP-BEZ235.

Table 2. PI3K Pathway Inhibitors in Clinical Trials

Targets	Compound (Company)	Study Population	Reported Efficacy/Responses	Reported Toxicities	References
PI3K/mTOR	SF-1126 (Semafore Pharmaceuticas)	Phase I: advanced solid tumors	SD in 11 of 28 pts	DLT: grade 3 diarrhea (one pt)	Chiorean et al ⁸²
	NVP-BEZ235 (Novartis)	Phase I/II: advanced solid tumors (breast cancer-enriched)	N/A	N/A	ClinicalTrials.gov
	NVP-BGT226 (Novartis)	Phase I/II: advanced solid tumors (including breast cancer)	N/A	N/A	ClinicalTrials.gov
	XL765 (Exelixis)	Phase I: refractory solid tumors	SD in five of 36 pts	Most frequent (> 10%) AEs: elevated liver enzymes, nausea, diarrhea Other AEs: anorexia/hypophosphatemia, rash, vomiting, and neurologic complaints	LoRusso et al ⁸³
PI3K	PX-866 (Oncothyreon)	Phase I: advanced solid tumors	SD in two of six pts (initial dosing cohorts)	Possible AEs: abdominal discomfort, mild diarrhea No DLTs in early dosing cohorts	Jimeno et al ⁸⁴
	XL147 (Exelixis)	Phase I: advanced solid tumors	SD (> 6 months) in six of 39 pts; one pt with HR CaP with normalized PSA	AEs: grade 3 rash (DLT in two of three pts at highest tested dose); grade 3 arterial thrombosis (one pt); grade 2 transaminitis (one pt); grade 1 hyperglycemia (four pts)	Shapiro et al ⁸⁵
	NVP-BKM120 (Novartis)	Phase I: solid tumors	N/A	N/A	Markman et al ⁸⁶
	GDC-0941 (Genentech/Piramed)	Phase I: advanced solid tumors	Evidence for potential antitumor activity in three of 19 pts	Most frequent AEs: Grade 1 to 2 nausea, fatigue, diarrhea, dysgeusia, and peripheral edema	Wagner et al ⁸⁷
	CAL-101 (Calistoga Pharmaceuticals)	Phase I: relapsed/refractory CLL or B-cell NHLs	PR in two of six pts, SD in four of six pts in early dosing cohorts	No AEs grade > 1 in first two dosing cohorts	Flinn et al ⁸⁸
Akt (allosteric)	MK-2206 (Merck)	Phase I: advanced solid tumors	SD in six of 19 patients	Most frequent AEs: skin (47.1%) and GI (41.2%) toxicities; DLTs: grade 3 to 4 rash, grade 3 mucositis	Tolcher et al ⁸⁹

Abbreviations: SD, stable disease; DLT, dose-limiting toxicity; Pt, patient; N/A, results from patient treatment in clinical trials have not been reported to date; AE, adverse event; HR CaP, hormone-refractory prostate cancer; PSA, prostate-specific antigen; CLL, chronic lymphocytic leukemia; NHL, non-Hodgkin's lymphoma; PR, partial response.

NVP-BEZ235 slowed the growth of PTEN-deficient human cancer cell line xenografts in mice, and it was well tolerated with no significant changes in body weight.⁹² Additionally, breast cancer cell lines with *HER2* amplification and/or *PIK3CA* mutations appeared to be particularly sensitive to these agents; however, it should be noted that only tumor stasis, and not tumor regression, was observed in vivo.⁹³ NVP-BEZ235 was further shown to induce apoptosis in estrogen-deprived estrogen receptor–positive breast cancer cells harboring either *PIK3CA* mutations or *PIK3CB* amplification.⁹⁴ In this study, RNA interference was also used to knock down p110 α , p110 β , or both in these cells. In some estrogen receptor–positive breast cancer cell lines harboring a *PIK3CA* mutation, dual knockdown of both p110 α and p110 β led to greater apoptosis following estrogen deprivation compared with knockdown of either isoform alone.⁹⁴ These results underscore a potential benefit of inhibiting both p110 α and p110 β , even in cancers that harbor specific genetic activation of one isoform.⁹⁴ A study with genetically engineered mice also demonstrated that NVP-BEZ235 was highly effective at shrinking murine lung tumors driven by a p110 α H1047R transgene.⁶⁵ Recently, phase I results for the dual PI3K-mTOR inhibitor, XL765 were reported at the 45th American Society of Clinical Oncology (ASCO) annual meeting (2009). There were no responses, but stable disease was noted in five of 36 patients.⁸³ There was evidence of 50% to 80% pathway inhibition in surrogate tissue. However, it is unclear whether this level of inhibition will be sufficient to induce shrinkage in potentially responsive tumors, or whether more complete inhibition will be required. No significant changes in serum glucose were noted, although an augmentation in food-induced plasma insulin increases was observed.⁸³

PI3K Inhibitors

The PI3K inhibitors can be divided into isoform-specific inhibitors or pan-PI3K inhibitors. Pan-PI3K inhibitors target all class I $_A$ PI3K in the cancer. These include wortmannin derivatives such as PX-886 or wortmannin prodrugs such as the self-activating viridans modified by dextran linker moieties that are designed to increase permeability and extend serum half-life.^{95–97} These agents exhibit cytostatic antitumor effects in vivo.^{95–97} The presence of *PIK3CA* mutations appears to predict for sensitivity to PX-886 across an array of cancer cell lines derived from different tissues of origin.⁷⁷ Interestingly, PTEN loss also appears to predict for PX-886 sensitivity, despite its relatively low efficacy toward p110 β .⁷⁷ Animals treated with PX-886 experienced hyperglycemia with decreased glucose tolerance as a major toxicity of PI3K inhibition, but this could be overcome with the oral antidiabetic agent pioglitazone.⁹⁸ Phase I clinical trial results for other pan-PI3K inhibitors have also been reported. Of 19 patients with solid tumors treated with GDC-0941 (Piramed/Genentech, Slough, United Kingdom/South San Francisco, CA), three demonstrated potential signs of antitumor activity.⁸⁷ Another pan-PI3K inhibitor, XL147, produced durable disease control in six of 39 treated patients. As observed with the dual PI3K-mTOR inhibitor XL765, plasma glucose levels were minimally affected by XL147, although an augmentation of food-induced plasma insulin increases was noted.⁸⁵

A possible advantage of isoform-specific PI3K inhibitors is that they may be tolerated at doses resulting in more complete target inhibition with fewer adverse effects. Isoform-specific inhibitors that selectively inhibit p110 α , β , δ , or γ catalytic subunits are under investigation in preclinical studies.^{99,100} Indeed, a p110 δ -specific inhibitor tested for refractory non-Hodgkin's lymphoma and chronic lympho-

cytic leukemia induced responses in 6 out of 12 patients (presented at the 45th Annual Meeting of ASCO in 2009).⁸⁰

AKT Inhibitors

Both adenosine triphosphate (ATP) mimetics and noncatalytic-site AKT inhibitors are under active clinical development.^{90,101} Cancers with *AKT1* mutations and *AKT1* and *AKT2* amplifications may be expected to be among the more sensitive to AKT inhibitors. However, this class of inhibitors will not block the non-AKT effectors of PI3K signaling and, paradoxically, could actually increase PI3K-dependent activation of those effectors via loss of negative feedbacks. This is especially important in light of the recent findings that the PDK1 substrate SGK3, and not AKT, may play a more prominent role in promoting PI3K-dependent viability in some cancers harboring *PIK3CA* mutations.¹⁰² Despite these findings, a recent study demonstrated that a noncatalytic site AKT1/AKT2 inhibitor was effective against breast cancer cell lines with *PIK3CA* mutations and *HER2* amplifications.¹⁰¹ Phase I results for the allosteric pan-AKT inhibitor MK-2206 (Merck, Whitehouse Station, NJ) showed stable disease in six of 19 patients and decreases in CA125 in patients with ovarian cancer. Adverse effects included rash and hyperglycemia.⁸⁹

mTOR Catalytic Site Inhibitors

Rapamycin interacts with FKBP12 in mammalian cells to form a complex that directly binds to the FKBP12-rapamycin-binding domain of mTOR in mTORC1, but not in mTORC2.^{103,104} Conversely, ATP-competitive mTOR inhibitors target the kinase domain of mTOR to impede the activity of both mTORC1 and mTORC2. Inhibiting mTORC2 would provide the theoretical advantage of blocking AKT activation. An ATP-competitive mTOR inhibitor might be more effective than rapamycin because, by blocking AKT activation, it would mitigate the activation of PI3K that often accompanies mTORC1 inhibition (ie, de-repression of the negative feedback, Fig 1). Intriguing preclinical data are emerging from studies of these compounds that shed new light on the potential limitations of rapamycin analogs. Feldman et al¹⁰⁵ demonstrated that two mTOR kinase domain inhibitors, PP242 and PP30, inhibit both mTORC1 and mTORC2. Unlike acute rapamycin treatment, which activates AKT, PP242 administration to mice suppressed AKT activation in tissues. PP242 was also a more effective inhibitor of proliferation than rapamycin.¹⁰⁵ Surprisingly, the improved efficacy of PP242 appeared to be due to more effective mTORC1 inhibition, rather than through its additional inhibition of mTORC2.¹⁰⁵ Similarly, the ATP-competitive mTOR inhibitor Torin1 impeded cell proliferation predominantly via its effects on mTORC1, not mTORC2.¹⁰⁶ Both PP242 and Torin1 were more effective inhibitors of 4E-BP1 phosphorylation and cap-dependent RNA translation than rapamycin.^{103,105,106} Three additional ATP-competitive mTOR inhibitors—WAY-600, WYE-687, and WYE-354—have been shown to inhibit proliferation of a variety of cancer cell lines more effectively than rapamycin, causing G1 cell cycle arrest, and in some cases, apoptosis.¹⁰⁴

Although the clinical results with PI3K pathway inhibitors are preliminary, their efficacy has modest at best. For their effective development, it will be imperative to understand why these drugs fail to produce a response when they do. Is the lack of activity due to inadequate inhibition of the target, or because complete inhibition of the target is not sufficient to produce antitumor activity? Indeed, most of the studies to date have not assessed this issue systematically. To

answer this question, future studies with quantitative pharmacodynamic assessments will be required to determine the degree of target inhibition. For example, a study with even a small number of patients with favorable genotypes (eg, *PIK3CA* mutants or *HER2* amplified) that correlates pharmacodynamic responses of PI3K pathway inhibition with outcomes may prove invaluable in identifying the reasons for lack of efficacy.

POTENTIAL CLINICAL USES FOR PI3K PATHWAY INHIBITORS

Thus far, preclinical studies have shown that PI3K pathway inhibitors may have significant single-agent activity in a few types of genetically defined cancers: *HER2*-amplified breast cancers, cancers with *PIK3CA* mutations, and *PTEN*-deficient cancers.^{65,77,92,93,101} To this point, data suggest that cancers with *KRAS* mutations may be fairly resistant to PI3K pathway inhibitors.^{65,77} Consequently, it seems likely that the presence of *KRAS* mutations will limit the efficacy of single-agent PI3K pathway inhibitors in cancers harboring both *KRAS* and *PIK3CA* mutations, such as many colon cancers.

In addition to these genetically defined settings, there may be other opportunities to target the PI3K pathway. For example, PI3K pathway inhibitors may be effective agents in the treatment of certain cancers that acquire resistance to RTK inhibitors. Cancers that are sensitive to receptor tyrosine kinase inhibitors (TKIs) have PI3K under the exclusive control of that RTK.⁶⁷ When a TKI works, it leads to downregulation of PI3K activity. For example in *HER2*-amplified breast cancers, trastuzumab disrupts the interaction between ErbB2 and ErbB3, resulting in ErbB3 dephosphorylation and loss of interaction with PI3K.¹⁰⁷ Furthermore, the presence of an activating *PIK3CA* mutation or depletion of *PTEN* correlates with a poor response to trastuzumab, presumably because these cancers fail to downregulate PI3K signaling in response to the anti-*HER2* therapy. Furthermore, when cancers that were initially sensitive to TKIs subsequently develop resistance, they invariably find a way to maintain PI3K signaling in the presence of the TKI.⁶⁷ These cancers therefore may be susceptible to addition of a PI3K pathway inhibitor to the TKI to re-induce remissions. This is an attractive approach in *HER2*-amplified breast cancers, because they appear to be sensitive to PI3K pathway inhibitors even before they develop resistance to anti-*HER2*-based therapies. Indeed, the *HER2*-amplified breast cancer cell lines with *PIK3CA* mutations are resistant to trastuzumab, and this can be overcome with treatment with GDC-0941.^{44,107} Similarly, *PTEN* loss, or activating mutations in *PIK3CA*, confers resistance to lapatinib, which can be overcome by treatment with NVP-BEZ235.⁶⁸

Potential of Combining PI3K With MEK Pathway Inhibitors

When cancers are sensitive to TKIs (ie, oncogene-addicted to an RTK), the TKI usually leads to downregulation of PI3K and other pathways, including the MEK-MAPK pathway. Thus, it remains unclear whether single-agent PI3K pathway inhibitors will promote dramatic responses (comparable to gefitinib in *EGFR*-mutant lung cancers or imatinib in chronic myelogenous leukemia), even in sensitive cancers. Most models of cancers that are sensitive to single-agent PI3K pathway inhibitors have demonstrated tumor stasis in vivo rather than frank tumor regressions.^{91-93,95-97} Consequently, it may be necessary to combine PI3K pathway inhibitors with other agents to

induce dramatic responses. Furthermore, there may be cancers that show no response to single-agent PI3K pathway inhibitors that will respond to PI3K pathway inhibitors combined with other therapies. For example, inhibition of PI3K signaling with NVP-BEZ235 failed to shrink established *Kras* G12D-driven lung tumors.⁶⁵ However, combined PI3K and MAPK pathway inhibition by treatment with NVP-BEZ235 and the MEK inhibitor ARRY-142886 led to marked tumor regression in this *Kras* lung cancer model.⁶⁵ Similarly, combined PI3K and MEK inhibition was required to effectively shrink *EGFR*-mutant lung cancers in genetically engineered mouse models.¹⁰⁸ Findings such as these are spurring the biotechnology and pharmaceutical industries to combine therapeutic inhibitors of these two pathways.

SUMMARY

Great strides are being made in our understanding of the diverse roles that PI3K signaling plays in cancer initiation, progression, and maintenance. Novel therapeutics targeting different components of this pathway are demonstrating efficacy in an array of human cancer types in preclinical studies, and these drugs are being carried forward into clinical trials. There is growing preclinical evidence that some genetically defined cancer subtypes may be the most sensitive to single-agent PI3K pathway inhibitors. These include cancers with *PIK3CA* activating mutations, loss of *PTEN*, and breast cancers with *HER2* amplification. However, it remains to be determined whether these sensitive cancers will demonstrate stable disease or tumor shrinkage in response to single-agent therapeutics. Conversely, cancers harboring activated Ras mutants appear to be insensitive to PI3K pathway inhibition alone. In such cases, effective treatment with PI3K inhibitors may require concomitant therapies targeting MAPK signaling.

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