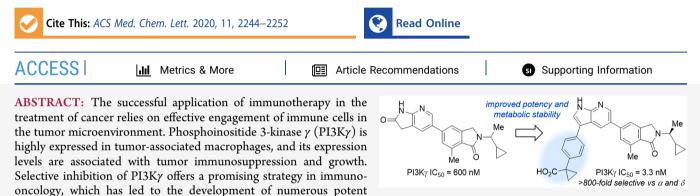
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Discovery of Potent and Selective 7-Azaindole Isoindolinone-Based PI3Kγ Inhibitors

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PI3K γ inhibitors with variable selectivity profiles. To facilitate further investigation of the therapeutic potential of PI3K γ inhibition, we required a potent and PI3K γ -selective tool compound with sufficient metabolic stability for use in future *in vivo* studies. Herein, we describe some of our efforts to realize this goal through the systematic study of SARs within a series of 7-azaindole-based PI3K γ inhibitors. The large volume of data generated from this study helped guide our subsequent lead optimization efforts and will inform further development of PI3K γ -selective inhibitors for use in immunomodulation.

KEYWORDS: PI3Ky, inhibitor, selective, azaindole, immunomodulation, cancer

P hosphoinositide 3-kinases (PI3Ks) are a class of enzymes responsible for the regulation of a wide variety of cellular activities such as signaling, survival, metabolism, and transport. A common substrate of these kinases is the glycerophospholipid phosphatidylinositol (PIP): PI3K-mediated phosphorylation of PIP initiates a signaling cascade via downstream proteins (e.g., protein kinase B (PKB), commonly referred to as Akt, and 3-phosphoinositide-dependent protein kinase-1 (PDPK1)) and ultimately mediates critical cellular functions, such as motility and proliferation.^{1,2} The class I PI3Ks, including PI3K α , PI3K β , PI3K γ , and PI3K δ , phosphorylate the C3-hydroxyl group of the inositol ring in phosphatidylinositol 4,5-bisphosphate.³

Although PI3Ks are involved in a variety of functions and are expressed throughout the body, PI3K γ and PI3K δ are of particular interest due to their expression in leukocytes as effectors of immune responses.^{4–7} In oncology, abnormalities in PI3K expression have been linked to cancer development, corresponding to high levels of PI3K γ expression and activity within immunosuppressive myeloid cells.^{8–18} Due to the significance of PI3K γ and PI3K δ in immunomodulation, many inhibitors with varying degrees of selectivity have been developed for the treatment of cancers and other pathologies of the immune system.^{19,20} Structures of several potent and γ -selective inhibitors have been published, selected examples of

which are shown in Figure 1a.²¹⁻²⁷ Among these, 1 (IPI-549, developed by Infinity Pharmaceuticals) is the only γ -selective inhibitor currently in clinical trials (Ph. 1/2).²⁸⁻³² Aryl sulfonamide 2 (developed by Exelixis), an advanced lead compound that resulted from optimization of a highthroughput screening hit, exhibits moderate PI3K isoform selectivity but also inhibits the mammalian target of rapamycin (mTOR) and other off-target kinases.²⁵ Recently, researchers at AstraZeneca reported a series of aminothiazole-based PI3K γ inhibitors such as 3, which was characterized by X-ray crystallography in complex with mouse PI3K δ (PDB ID The high isoform selectivity of 3 and its analogues 6FTN).² was attributed to the ability of the cyclopropylethyl group to displace the DFG motif of PI3K γ , which causes a unique γ isoform-specific conformational change.²⁷

As shown in Figure 1b using aryl sulfonamide **2** as a prototypical example, three regions exist within the ATP binding pocket of PI3K γ , which are typically considered during

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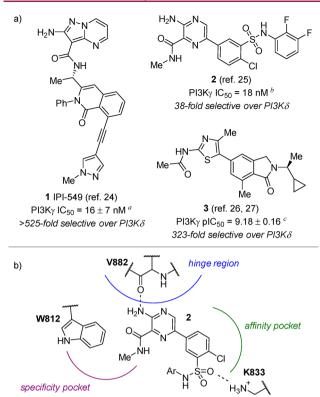


Figure 1. (a) Biochemical potency and selectivity values of PI3K γ -selective inhibitors, as reported in the literature (see Supporting Information for a comparison of literature and in-house assay protocols). Inhibition determined using the ADP-Glo assay (Promega). Reported ATP $K_m = 29 \pm 3.7 \ \mu$ M.³⁴ ^aAssay conditions: 3 mM ATP, 500 μ M dioctanoyl-phosphatidylinositol-4,5-bisphosphate (diC₈PIP₂) as substrate, and 40 nM PI3K γ . ^bAssay conditions: 1 μ M ATP, 10 μ M PIP₂, and 30 nM PI3K γ . ^cAssay conditions: 20 μ M ATP, 80 μ M diC₈PIP₂, and 1.2 nM PI3K γ . (b) Schematic of typical inhibitor binding modes within PI3K γ , exemplified by 2.²⁵

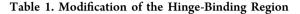
inhibitor design.³³ Interaction with the hinge region is considered essential for inhibitor activity; most known inhibitors engage in at least one hydrogen bond to V882. Inhibitor binding interactions within the affinity pocket are known to considerably influence both potency and selectivity against the other PI3K isoforms, as communicated in the discovery of 3.^{26,27} Most inhibitors that bind within the affinity pocket possess a functional group that engages K833 via hydrogen bonding or polar interactions. Finally, inhibitor binding within the specificity pocket also influences both potency and selectivity, as demonstrated in the discovery of 1.24 The W812 side chain in this region of the enzymeinhibitor complex is well-positioned for an edge-to-face interaction with the inhibitor's planar motif, as confirmed by the recently reported X-ray co-crystal structure of 1 in complex with human PI3Ky (PDB ID 6XRL).³⁵

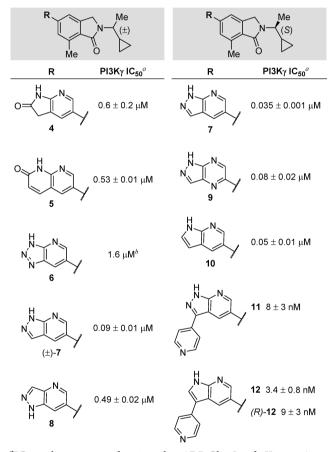
Considering the promise of selective PI3K γ inhibition for cancer immunotherapy, we pursued the design of novel inhibitor structures, utilizing the extensive body of medicinal chemistry knowledge as a source of inspiration. Our primary objective was to design a molecule possessing (1) similar or greater PI3K γ inhibitory potency and isoform selectivity compared to known inhibitors, such as 1, as measured in both biochemical and cell-based PI3K γ inhibition assays, and (2) suitable ADME properties for use in future *in vivo* studies. A molecule fulfilling these criteria would enable the eventual

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deconvolution and appreciation of the role of PI3K γ inhibition in the context of our broader immunomodulatory work. To realize these goals, we embarked upon the discovery of novel inhibitors, initially through a pharmacophore mapping approach based on visual comparison of known literature inhibitors followed by a systematic SAR investigation of promising lead molecules, some of which have recently been described.³⁵

Due to the high isoform selectivity and potency of 3 (IC₅₀ = 4 ± 1 nM, as determined in our in-house biochemical assays, which measure a compound's ability to inhibit the kinase activity of PI3K; see Supporting Information for details) and its analogues,^{26,27} we designed an initial set of compounds by attempting to mimic the hinge-binding acylaminothiazole moiety with a variety of functionally similar heterocycles (Table 1). Most of the compounds investigated contained both

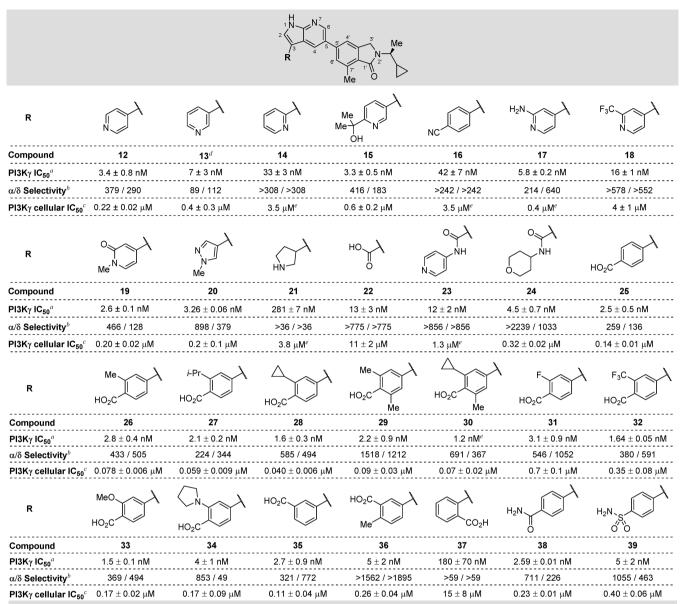




 a IC₅₀ values measured using the ADP-Glo Lipid Kinase Assay (Promega) with 25 μ M ATP and 50 μ M phosphatidylinositol 4,5-bisphosphate as substrate. b Compound was tested once.

a hydrogen bond donor and a hydrogen bond acceptor motif, which could potentially create a stronger interaction with V882. We initiated our studies with racemic compounds 4 and 5, which bear a structural resemblance to 3. Although potency was substantially reduced for these compounds ($IC_{50} = 0.6$ and $0.53 \ \mu$ M, respectively), we were encouraged by the fact that replacement of the aminothiazole motif of 3 did not eliminate activity. Although triazolopyridine **6** showed decreased potency ($IC_{50} = 1.6 \ \mu$ M), a less acidic analogue, (\pm)-7, was significantly more active ($IC_{50} = 0.09 \ \mu$ M). In contrast, an

Table 2. Evaluation of Azaindole C3 Substituents



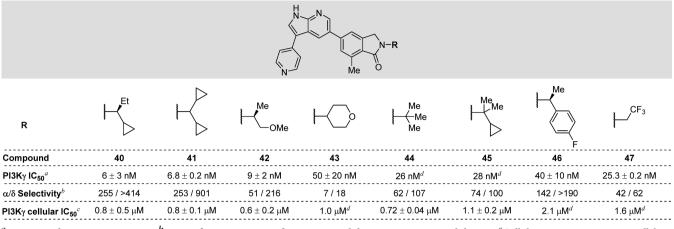
^{*a*}Enzymatic kinase activity assay. ^{*b*}Ratio of PI3K α or PI3K δ enzymatic inhibition over PI3K γ inhibition. ^{*c*}Determined using the AlphaLISA SureFire Ultra Akt assay (PerkinElmer) with THP-1 cells. ^{*d*}Racemate. ^{*e*}Compound was tested once.

isomeric analogue, **8**, in which the pyrazole NH group is not properly positioned for interaction with V882, was significantly less potent (IC₅₀ = 0.49 μ M), confirming the importance of a hydrogen bond donor and acceptor pair for interaction with the hinge region. The stereochemical configuration of these compounds was also shown to be important, as 7, the (*S*) enantiomer of (±)-7, showed improved activity (IC₅₀ = 0.035 μ M) compared to the racemic compound. Although enantiopure pyrazolopyrazine **9** showed a 2-fold reduction in potency (IC₅₀ = 0.08 μ M) relative to 7, azaindole **10** (IC₅₀ = 0.05 μ M) had similar inhibitory activity to 7.

Based on these results and earlier studies by the Meng and Zhou laboratories,³⁶ we evaluated the incorporation of a 4-pyridyl moiety at the C3 position of the two most promising scaffolds (i.e., azaindazole 7 and azaindole 10), which we envisioned would provide favorable interactions with the specificity pocket of PI3K γ . Consistent with our hypothesis,

azaindazole 11 (IC₅₀ = 8 nM) was 4-fold more potent than 7, while azaindole 12 (IC₅₀ = 3.4 nM) showed an even greater improvement in potency (approximately 15-fold) relative to the unsubstituted analogue, 10. (*R*)-12 was 3-fold less potent than 12, consistent with the previously reported superiority of the (*S*) enantiomer of the isoindolinone moiety of 3.^{26,27} These initial studies established 12 as a suitable starting point for further optimization.

Our subsequent efforts centered on evaluation of various C3-substituted derivatives of **12** (Table 2). To better distinguish between compounds that generally displayed low-nanomolar potency in the biochemical kinase activity assay, we further characterized these inhibitors under more physiologically relevant conditions using a functional cellular inhibition assay, which measures the extent of PI3K-mediated phosphorylation of Akt. To assess PI3K isoform selectivity, we employed routine enzymatic kinase activity assays for the α and δ PI3K Table 3. Modification of the N2' Substituent



^{*a*}Enzymatic kinase activity assay. ^{*b*}Ratio of PI3K α or PI3K δ enzymatic inhibition over PI3K γ inhibition. ^{*c*}Cellular potency in THP-1 cell line. ^{*d*}Compound was tested once.

isoforms only, as most of the compounds in this series were inactive against the β isoform at concentrations up to 10 μ M (>1000-fold selectivity). We were pleased to find that compound 12 exhibited submicromolar activity in THP-1 cells (THP-1 IC₅₀ = 0.22 μ M), with excellent isoform selectivity (379-fold vs PI3K α and 290-fold vs PI3K δ). The observed difference between biochemical and cellular potency was not surprising, as cellular potency is frequently impacted by a compound's permeability, protein binding, and off-target activity as well as differences in the concentration of ATP present under the assay conditions.

Although the 3-pyridyl analogue 13 ($IC_{50} = 7$ nM, THP-1 $IC_{50} = 0.4 \ \mu M$) showed comparable potency to 12, the 2pyridyl analogue (14) was substantially less active ($IC_{50} = 33$ nM, THP-1 IC₅₀ = 3.5 μ M), indicating a hydrogen bond acceptor extending outward from the core is advantageous.³ Other modifications preserving this paradigm (e.g., 15-19) were either tolerated or had reduced activity, with biochemical potency values ranging from 2.6 to 42 nM, and THP-1 cell potencies ranging from 0.20 to 3.5 μ M. Pyrazole 20 (IC₅₀ = 3.26 nM, THP-1 IC₅₀ = 0.2 μ M) showed comparable potency to 12, but incorporation of a saturated, basic amine substituent (e.g., 21, IC₅₀ = 281 nM, THP-1 IC₅₀ = 3.8μ M) proved to be detrimental to activity. Interestingly, carboxylic acid 22 and its amide derivatives, 23 and 24, were moderately active in the biochemical assay (IC₅₀ = 13, 12, and 4.5 nM, respectively) despite being structurally distinct, although cellular potencies were more variable (THP-1 IC₅₀ = 11, 1.3, and 0.32 μ M, respectively).

In addition to 5- and 6-membered aromatic heterocycles, benzoic acids were identified as suitable C3 substituents. *para*-Benzoic acid **25** (IC₅₀ = 2.5 nM, THP-1 IC₅₀ = 0.14 μ M), as well as several substituted benzoic acid analogues (**26**–**30** and **33**–**35**), exhibited improved cellular potency compared to **20**. The *ortho*-methyl, -isopropyl, and -cyclopropyl analogues of **25** showed an encouraging increase in cellular potency (THP-1 IC₅₀ = 0.078, 0.059, and 0.040 μ M, respectively). Disubstituted analogues of **25** (e.g., **29** and **30**, THP-1 IC₅₀ = 0.09 and 0.07 μ M, respectively), for which steric effects limit the conformational freedom of the carboxylate group and may attenuate productive interactions within the affinity pocket, showed a minimal reduction in potency relative to the monosubstituted counterparts. Electron-poor benzoic acid derivatives (e.g., **31**

and **32**, THP-1 IC₅₀ = 0.7 and 0.35 μ M, respectively) showed a substantial decrease in potency, while analogues bearing electron-donating methoxy or pyrrolidine *ortho*-substituents (e.g., **33** and **34**, THP-1 IC₅₀ = 0.17 μ M for both) showed similar potency to unsubstituted benzoic acid **25**.

Interestingly, while *meta*-benzoic acid 35 (THP-1 IC_{50} = 0.11 μ M) exhibited similar potency to the *para*-analogue (25), introduction of an ortho-methyl substituent to 35 resulted in a 2-fold reduction in potency (36, THP-1 IC₅₀ = 0.26μ M). This observation further corroborates the potential effect of the conformation of the carboxylate group on compound potency. In contrast to 25 and 35, ortho-benzoic acid derivative 37 exhibited a dramatic reduction in potency (THP-1 $IC_{50} = 15$ μ M), consistent with the trends observed for isomeric pyridine analogues 12-14. Amide and sulfonamide analogues of 25 generally exhibited reduced potency (38 and 39, THP-1 IC_{50} = 0.23 and 0.40 μ M, respectively). For the compounds shown in Table 2, cellular potencies generally correlated with biochemical potency. Apart from compounds with a particularly unique geometry (21, 22, and 37), a modest correlation was observed between cellular potency and biochemical assayderived lipophilic efficiency values (see Supporting Information for details). Although there was substantial variation of isoform selectivity among the assayed compounds, most of the active compounds exhibited >100-fold selectivity against both PI3K α and δ . The highest isoform selectivity was observed for amide 24 and benzoic acid analogues 29 and 36.

During the course of these investigations into the SAR of the C3 substituent, we also concurrently investigated the SAR of the N2' isoindolinone substituent of 12 (selected due to its combination of potency, selectivity, and structural simplicity), which we envisioned would be situated within the affinity pocket of the PI3Ky-inhibitor complex (Table 3). Replacement of the methyl group of the cyclopropylethyl moiety with either an ethyl group (40, IC₅₀ = 6 nM, THP-1 IC₅₀ = 0.8 μ M) or a cyclopropyl group (41, $IC_{50} = 6.8$ nM, THP-1 $IC_{50} = 0.8$ μ M) resulted in similar biochemical potency and selectivity but approximately 4-fold reduction in cellular potency. On the other hand, replacement of the cyclopropyl group with a methoxymethylene group (see 42, $IC_{50} = 9 \text{ nM}$, THP-1 $IC_{50} =$ $0.6 \,\mu\text{M}$) resulted in a 3-fold reduction in both biochemical and cellular potency and a substantial reduction in isoform selectivity, confirming the important contribution of the

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R	 Me	 <i>i</i> -Pr	 OMe	CI	CN	CF3			
Compound	22	48	49	50	51	52			
ΡΙ3Κγ ΙC ₅₀ ^α	$3.26\pm0.06~\text{nM}$	4.2 ± 0.8 nM	$4.3\pm0.6~\text{nM}$	$2.4\pm0.2~\text{nM}$	$3.7\pm0.2~\text{nM}$	$2.4\pm0.9~\text{nM}$			
x/δ Selectivity ^b	898 / 379	>1666 / 1378	455 / 365	1059 / 533	2795 / 244	1049 / 266			
PI3Kγ cellular IC ₅₀ °	0.2 ± 0.1 μM	$0.3\pm0.1~\mu M$	$0.17\pm0.07~\mu\text{M}$	$0.17\pm0.09~\mu M$	$0.17\pm0.05~\mu\text{M}$	$0.09\pm0.02~\mu\text{M}$			
R	0 NHMe	NHAc	o=s ó́NHMe	O=S o″NMe₂	 NHMs	O=S ÓMe			
Compound	53	54	55	56	57	58			
ΡΙ3Κγ ΙC ₅₀ ^{<i>a</i>}	$7.9\pm0.5~\text{nM}$	$2.5\pm0.9~\text{nM}$	4 ± 1 nM	$2.6\pm0.9~\text{nM}$	4 ± 3 nM	1.5 ± 0.3 nM			
u/δ Selectivity ^b	>1060 / >1183	>4159 / >2743	254 / 101	449 / 159	1022 / 615	642 / 289			
PI3Kγ cellular IC ₅₀ °	$0.73\pm0.08~\mu M$	0.5 ± 0.3 μM	$0.07\pm0.01~\mu M$	$0.10 \pm 0.01 \ \mu M$	$0.15 \pm 0.01 \ \mu M$	0.05 ± 0.02 μN			

Table 4. Modification of the C7' Substituent of 20

cyclopropyl group to isoform selectivity. An even greater reduction in potency and selectivity was observed for tetrahydropyran 43 (IC₅₀ = 50 nM, THP-1 IC₅₀ = 1.0 μ M), which reinforces the importance of the specific size and shape of the N-alkyl substituent. Analogues bearing a sterically bulky *N*-alkyl substituent, such as a *t*-butyl group (44, $IC_{50} = 26$ nM, THP-1 IC₅₀ = 0.72 μ M), showed diminished potency and isoform selectivity. Similarly, reduced potency and selectivity were observed for analogue 45 (IC₅₀ = 28 nM), which bears an additional methyl group relative to 12 and is achiral. Despite its large size, arylethane 46 retained modest activity in the biochemical assay (IC₅₀ = 40 nM), but its potency in cells was significantly decreased (THP-1 IC₅₀ = 2.1 μ M) relative to 12. Compound 47 (THP-1 IC₅₀ = 1.6 μ M), which bears an unbranched trifluoroethyl group, exhibited a significant reduction in both potency and selectivity compared to 12. Taken together, these results indicated that proper substitution of the isoindolinone nitrogen is critical to achieving high potency and selectivity. Moreover, these studies further supported the importance of the cyclopropylethyl group to achieving the optimal balance of potency and isoform selectivity exemplified by 12.

Further characterization of **12** revealed that it was a potent inhibitor of the major CYP isoforms (IC₅₀ values for the 1A2, 2D6, 2C9, 2C19, and 3A4 CYP isoforms were 0.7, 2.0, 0.3, <0.1, and 0.3 μ M, respectively), possibly due to pyridine—iron coordination. On the other hand, pyrazole **20**, which lacks the pyridine moiety, showed an improved CYP inhibition profile (IC₅₀ values for the 1A2, 2D6, 2C9, 2C19, and 3A4 CYP isoforms were 2.9, 26, 1.4, 0.2, and 0.8 μ M, respectively). Based on its decreased CYP inhibition and comparable cellular potency to **12**, **20** was selected for further exploration of the isoindolinone substituent.

As shown in Table 4, improvement of cellular potency could be realized while maintaining isoform selectivity through various steric and electronic modifications of the C7' group of **20**. Incorporation of a sterically bulky isopropyl group (**48**, THP-1 IC₅₀ = 0.3 μ M) had little impact on potency or selectivity. Similarly, incorporation of an electron-donating methoxy group (49, THP-1 IC₅₀ = 0.17 μ M) or electronwithdrawing chloro- or nitrile groups (e.g., **50** and **51**, THP-1 IC₅₀ = 0.17 μ M for both) also had little effect on potency or selectivity. Importantly, trifluoromethyl analogue **52** (THP-1 IC₅₀ = 0.09 μ M exhibited a 2-fold improvement in potency over **20** while maintaining high isoform selectivity. While both acetamide isomers **53** and **54** had reduced activity (THP-1 IC₅₀ = 0.73 and 0.5 μ M, respectively) relative to **20**, C7'-sulfonamide and -sulfone analogues exhibited up to a 4-fold improvement in potency (**55-58**, THP-1 IC₅₀ = 0.05-0.15 μ M). Despite the promising potency of **55** and **58**, these compounds exhibited extrahepatic clearance in rat models (6.0 and 4.5 L/h/kg, respectively), which precluded their further progression.

Faced with the CYP inhibition liabilities of C3-pyridine analogues and the poor metabolic stability of the C3-pyrazole series, we turned our attention to the C3-benzoic acid series, exemplified by analogues 25, 26, and 29 (Table 5). Benzoic acid 25 and the corresponding ortho-substituted analogue, 26, both exhibited poor in vitro metabolic stability, as measured in human and rat hepatocytes, with rat intrinsic clearance (CL_{int}) values $\geq 40 \ \mu L/min/10^6$ cells. While hepatocyte stability could be improved through the incorporation of an additional ortho substituent on the benzoic acid moiety (29, rat CL_{int} 2.0 $\mu L/$ $min/10^{6}$ cells), this compound exhibited extrahepatic clearance in rat models (e.g., CL = 6.9 L/h/kg). While the C7'-CF₃ analogues of 26 and 29 (see 59 and 60, THP-1 $IC_{50} = 0.04$ and 0.03 μ M, respectively) demonstrated improved potency consistent with our previous observations in the C3-pyrazole series-their metabolic stability was comparable to the C7'-Me analogues. The poor in vitro/in vivo correlation for this series of compounds may be due to active efflux, as Caco-2 permeability studies revealed parent compound 25 to have low permeability ($P_{app} = 1.07 \times 10^{-6} \text{ cm/sec}$) and an efflux ratio of 46.6.

Finally, a breakthrough came with the synthesis of C3 phenylacetic acid derivatives **61** (THP-1 IC₅₀ = $0.16 \,\mu$ M) and

Table 5. Further Characterization and SAR of C3 Benzoic Acid and Phenylacetic Acid Azaindole Derivatives

		R_1	R_2 N Me		
R ₁	HO ₂ C	Me HO ₂ C	Me HO ₂ C	Me HO ₂ C Me	Me HO ₂ C Me
Compound	25 (R ₂ = Me)	26 (R ₂ = Me)	59 (R ₂ = CF ₃)	29 (R ₂ = Me)	60 (R ₂ = CF ₃)
ΡΙ3Κγ ΙC ₅₀ ^α	$2.5\pm0.5~\text{nM}$	$2.8\pm0.4~\text{nM}$	1.2 ± 0.1 nM	$2.2\pm0.9~\text{nM}$	2.8 nM ^d
α/δ Selectivity ^b	259 / 136	433 / 505	918 / 335	1518 / 1212	257 / 84
PI3Kγ cellular IC ₅₀ ^{c}	$0.14\pm0.01~\mu\text{M}$	$0.08\pm0.01~\mu\text{M}$	$0.04\pm0.01~\mu\text{M}$	$0.09\pm0.03~\mu\text{M}$	$0.03\pm0.01~\mu\textrm{M}$
Hepatocyte CL _{int} (μL/min/10 ⁶ cells)	human: 24 rat: 54	human: 19 rat: 51	human: 20 rat: 46	human: <1.2 rat: 2.0	human: 1.8 rat: <1.2
<i>in vivo</i> CL (rat) ^e	11 L/h/kg	n.d.	n.d.	6.9 L/h/kg	28 L/h/kg
R ₁	CO ₂ H	Me CO ₂ H	Me Me CO ₂ H	CO ₂ H	CO ₂ H
Compound	61 (R ₂ = Me)	62 (R ₂ = Me)	63 (R ₂ = Me)	64 (R ₂ = Me)	65 (R ₂ = CF ₃)
ΡΙ3Κγ ΙC ₅₀ ^α	$2.6\pm0.2~\text{nM}$	3.3 nM ^d	2.1 nM ^d	$3.3\pm0.4~\text{nM}$	$3.6\pm0.6~\text{nM}$
α/δ Selectivity ^b	878 / 872	539 / 397	835 / 968	863 / 1401	742 / 620
PI3Kγ cellular IC ₅₀ c	$0.16\pm0.02~\mu M$	$0.15\pm0.07~\mu M$	$0.18\pm0.02~\mu\text{M}$	$0.15\pm0.04~\mu M$	$0.11\pm0.02~\mu M$
Hepatocyte CL_{int} (µL/min/10 ⁶ cells)	human: 6.9 rat: 17	human: 8.0 rat: 23	human: 2.3 rat: 27	human: 1.7 rat: 16	human: 2.6 rat: 18
<i>in vivo</i> CL (rat) ^e	n.d.	1.8 L/h/kg	n.d.	1.8 L/h/kg	n.d.

^{*a*}Enzymatic kinase activity assay. ^{*b*}Ratio of PI3K α or PI3K δ enzymatic inhibition over PI3K γ inhibition. ^{*c*}Cellular potency in THP-1 cell line. ^{*d*}Compound was tested once. ^{*e*}n.d. = value not determined.

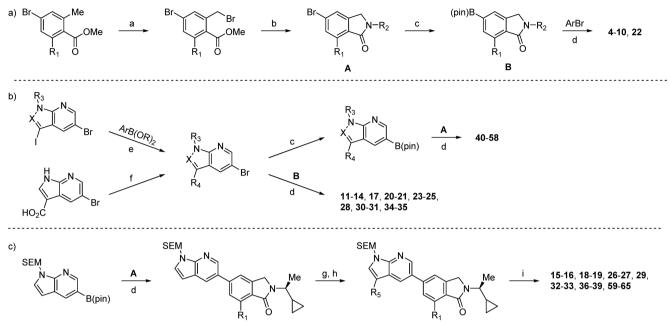
62 (THP-1 IC₅₀ = 0.15 μ M). Although these molecules showed somewhat diminished inhibitory activity relative to the benzoic acids, they retained high isoform selectivity and displayed an encouraging improvement in hepatocyte stability (rat $CL_{int} = 17$ and 23 $\mu L/min/10^6$ cells for 61 and 62, respectively). Moreover, compound 62 exhibited significantly lower in vivo clearance (CL = 1.8 L/h/kg) relative to the benzoic acid analogues, which lack a methylene spacer. Gratifyingly, substitution of the α -methylene group of 61 (see 63–65, THP-1 IC₅₀ = 0.11–0.18 μ M) resulted in further improvements in human hepatocyte stability without substantially altering the potency or selectivity of these inhibitors. While this optimization campaign did not ultimately deliver a compound suitable for further development, compound 64 emerged as the most promising inhibitor due to its high cellular potency, excellent isoform selectivity, and acceptable pharmacokinetic profile.

The general synthetic routes used to access our azaindole/ azaindazole inhibitors are shown in Scheme 1. The overall sequence is highly modular and allows for the introduction of diversity in several independent steps. Depending on the specific substrate, conditions such as the coupling partner, palladium catalyst, base, or solvent may vary (see Supporting Information for details). In general, the respective coupling steps can be done in any order to facilitate SAR exploration.

As shown in Scheme 1a, isoindolinone intermediates A and B are generated from methyl benzoate and primary amine precursors. These can be coupled to a variety of heteroaryl bromides to yield various simple inhibitors (4-10, 22). Alternatively, A or B can be coupled with C3-substituted azaindole/azaindazole intermediates, as shown in Scheme 1b. This sequence proved particularly useful during SAR evaluation of isoindolinone substituents. Finally, the sequence shown in Scheme 1c enabled rapid evaluation of C3 aryl substituents in combination with the most promising isoindolinone scaffolds.

Traditionally, ligand interactions within the affinity and specificity pockets of PI3K γ have been thought to independently affect potency and selectivity, respectively. In contrast, our SAR investigations demonstrate that improvements in both metrics can be realized through careful design of inhibitors that are capable of occupying both regions of the enzyme active site. Through systematic SAR analysis, we have discovered a series of novel, potent, and selective azaindole-based PI3K γ inhibitors. These compounds exhibit cellular IC₅₀ values as low as 0.040 μ M (**28**), while maintaining >300-fold selectivity against all other class I PI3K isoforms. In further

Scheme 1. General Synthetic Routes to Azaindole-Based Inhibitors^a



^aReagents and conditions: (a) NBS, (PhCO₂)₂, CCl₄, 80 °C; (b) RNH₂, B(OH)₃, K₂CO₃, ACN, r.t.; (c) B₂pin₂, (dppf)PdCl₂, KOAc, dioxane, 100 °C; (d) (dppf)PdCl₂, Na₂CO₃, dioxane/H₂O, 100 °C; then deprotection (conditions vary, see Supporting Information); (e) (dppf)PdCl₂, Na₂CO₃, dioxane/H₂O, 80 °C; (f) RNH₂, HATU, *i*-Pr₂NEt, DMF, 40 °C; (g) NBS, CH₂Cl₂, r.t.; (h) ArB(OR)₂, (dppf)PdCl₂, Na₂CO₃, dioxane/H₂O, 100 °C; (i) TFA, r.t.; *N*,N-DMEDA, MeOH, 45 °C.

inhibitor design iterations, C3 phenylacetic acid derivatives (e.g., **64**) exhibited significantly improved pharmacokinetic properties, helping pave the way for future biological studies. The volume of data generated in this study will help expedite future campaigns toward potent and orally bioavailable PI3K γ inhibitors for potential clinical applications.

ASSOCIATED CONTENT

Supporting Information

The Supporting Information is available free of charge at https://pubs.acs.org/doi/10.1021/acsmedchemlett.0c00387.

Biological assay procedures, synthetic procedures and characterization of PI3Ky inhibitors (PDF)

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Notes

The authors declare the following competing financial interest(s): All authors are current or former employees of Arcus Biosciences.

ABBREVIATIONS

SAR, structure—activity relationship; ATP, adenosine triphosphate; ADME, absorption, distribution, metabolism, and excretion; CYP, cytochrome P450; NBS, *N*-bromosuccinimide; ACN, acetonitrile; r.t., room temperature (23 °C); pin, pinacolato; dppf, 1,1'-bis(diphenylphosphino)ferrocene; HATU, (1-[bis(dimethylamino)methylene]-1*H*-1,2,3-triazolo-[4,5-*b*]pyridinium 3-oxide hexafluorophosphate; DMF, *N*,*N*dimethylformamide; SEM, 2-(trimethylsilyl)ethoxymethyl; TFA, trifluoroacetic acid; *N*,*N*-DMEDA, *N*,*N*-dimethylethane-1,2-diamine

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