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Letter

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Discovery of a Selective Phosphoinositide-3-Kinase (PI3K)-γ Inhibitor (IPI-549) as an Immuno-Oncology Clinical Candidate

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KEYWORDS: IPI-549, PI3K-gamma inhibitor, phosphoinositide-3-kinase, isoform selectivity, neutrophil migration, immuno-oncology

ABSTRACT: Optimization of isoquinolinone PI3K inhibitors led to the discovery of a potent inhibitor of PI3K- γ (**26** or IPI-549) with >100-fold selectivity over other lipid and protein kinases. IPI-549 demonstrates favorable pharmacokinetic properties and robust inhibition of PI3K- γ mediated neutrophil migration in vivo and is currently in Phase 1 clinical evaluation in subjects with advanced solid tumors.

Phosphoinositide-3-kinases (PI3Ks) belong to a family of signal transducing enzymes that mediate key cellular functions in cancer and immunity. The primary role of these lipid kinases entails catalysis of the phosphorylation of the inositol ring of membrane phosphoinositides, which then serve as sites for the activation of secondary messengers involved in the regulation of multiple biological processes, including cell growth, survival, differentiation, proliferation and migration. PI3Ks are divided into three classes (I, II, and III) based on substrate specificity, sequence homology, and types of regulatory subunits. Class I PI3Ks are further divided into two groups: Class IA, which includes PI3K- α , β , and δ , and Class IB, which contains only one member, PI3K-gamma (y). While Class IA PI3Ks are predominantly activated by receptor tyrosine kinase signaling, PI3K- γ is activated primarily by G-protein coupled receptors.1-6

The PI3K-y isoform is expressed in immune cells and has limited, if any, expression in normal or malignant epithelial cells and connective tissue cells.⁷ Studies of PI3K-y knockout mice show that PI3K-y is important for cellular activation and migration in response to certain chemokines.⁷⁻⁸ PI3K-y signaling is particularly important for the function of myeloid cells, where it is downstream of G-protein coupled receptors (GPCRs) (e.g., chemokine receptors) and RAS.⁷⁻⁹ In addition, PI3K- γ can be activated in response to tissue hypoxia in these cells.¹⁰ A critical role for PI3K-y in the unique myeloidderived cells that constitute a key component of the immune-suppressive tumor microenvironment has also been demonstrated in PI3K-y deletion and kinase-dead knock-in studies. For example, murine syngeneic tumors grow slower when transplanted into immune-competent mice where PI3K-y is genetically inactivated.^{9,10} This growth reduction is due to the abrogation of tumor-associated myeloid cells that are

known to promote an immune-suppressive tumor microenvironment that permits tumor growth.⁹⁻¹² In addition, tumor-associated myeloid cells are postulated to support tumor regrowth after radiation or chemotherapy, and to enable metastatic spread.¹³ These preclinical studies highlight a critical role for PI3K- γ in myeloid cell biology, and suggest that PI3K- γ inhibition in tumor-associated myeloid cells may be effective at preventing tumor growth in a variety of settings.

To test the hypothesis that pharmacological inhibition of PI3K- γ in tumor-associated myeloid cells could block their immune-suppressive function and lead to enhanced immune attack on tumor cells, a potent and selective synthetic small-molecule inhibitor of the PI3K- γ isoform was needed. Although inhibitors of PI3K- γ have been reported over the past decade, their selectivity for PI3K- γ over other PI3K isoforms or pharmacologic properties were until recently not suitable for assessment of inhibiting only PI3K- γ in vivo.¹⁴⁻²³ Hence, we set out to identify potent and selective inhibitors of PI3K- γ with desirable drug-like properties.

The *N*-phenyl 8-chloroisoquinolinone motif served as a possible scaffold to explore alternative hinge binding motifs that may provide unique interactions with PI3K- γ .²⁴ To this end, we assembled a focused collection of 8-chloroisoquinolinone analogs that could be readily synthesized having hinge binding motifs containing at least one hydrogen bond acceptor and 0-2 hydrogen bond donors. We screened this focused collection for their ability to inhibit PI3K- γ and PI3K- δ at physiological ATP concentrations (3 mM) and were pleasantly surprised to see that compounds 1 and 2 showed moderate selectivity for PI3K- δ (Table 1).

Table 1. Structure-activity relationship from hinge (R_1) and linker modifications (R_2) .



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Cmpd	R_1	R ₂	Biochemical IC ₅₀ $(nM)^a$		
			ΡΙ3Κ-γ	PI3K-δ	
1	N-N N-N	Me	40	400	
2	N NH ₂	Me	60	320	
3	N-N-N	Me	400	800	
4	N N N	Me	1000	900	
5	N-N	Me	300	2600	
6	NH ₂	Me	>10000	>10000	
7	N-N-N	Н	>7000	>10000	
8		Et	150	600	

^{*a*} Data reported as the average of at least two runs. See Supporting Information for description of assay conditions and SEM values.

The hinge binding motifs on compounds 1 and 2 shared the following common features: an amide bond to isoquinolinone core with a chiral linker, an exocyclic amine as part of a hydrogen bond donor/acceptor pair, and a heterocyclic nitrogen oriented towards the core. The importance of these structural features on potency and selectivity was confirmed by a set of direct analogs of 1 and 2 (Table 1). Analogs without the exocyclic amine (3 and 4) were significantly less potent and selective than 1 and 2, respectively. The carbo-analog of the 6-5 bicyclic system (5) was less active than the aza-analog (1), whereas the same modification on the 6-6 bicyclic system (6 vs 2) rendered the compound largely inactive against both PI3K isoforms. These data suggest that the dihedral angle between the carbonyl amide and the heterocyclic hinge binding motif 1 and 2 provides a preferred conformation for binding to both PI3K- γ and PI3K- δ . Changes to the chiral linker between the hinge binding motif and N-phenyl 8-chloroisoquinolinone core also proved to be deleterious to PI3K potency and selectivity. The enantiomer of 1 and the des-methyl achiral analog 7 were largely inactive against both PI3K isoforms. Interestingly, the selectivity was reduced when the chiral methyl is extended to an ethyl substituent (8 vs 1). These data highlight key hydrogen bonding features on the hinge binding motifs and the preferred chiral linker of *N*-phenyl 8chloroisoquinolinones 1 and 2 that make them bona fide leads for the development of selective PI3K- γ inhibitors.

We evaluated leads 1 and 2 for selectivity across other PI3K isoforms and found 1 to be remarkably selective across other Class I and Class II PI3K isoforms (Table S1). Therefore, we focused our attention on further optimization of compound 1. Taking the published co-crystal structures of propeller-shaped ligand PIK-39 with PI3K-8 and PI3K-y into consideration, we evaluated C8 substitution on compound 1 to access a nonconserved region within PI3Ks with the aim of improving the PI3K-γ potency and selectivity (Table 2).^{25, 26} We found small C8 substitutions such as methyl (9) and fluoro (10) did not have a profound impact on PI3K-y potency or selectivity. However, a significant loss in PI3K-y potency was seen with an electron-withdrawing cyano group (11) or electrondonating groups (12-14) at the 8 position. Vinyl substitution at the 8 position (15) also had a negative impact on PI3K- γ potency. However, 8-alkyne substitutions (16 and 17) unexpectedly provided modest improvements in potency for PI3K-y over the 8-chloro substitution in 1. Importantly, the methyl alkyne analog 17 also demonstrated weaker activity against PI3K- δ compared to 1 and 16, and thus 17 had >40-fold selectivity for PI3K- γ over PI3K- δ , suggesting that the alkyne substitution makes significant non-favorable interactions with PI3K-8 at the non-conserved residues adjacent to the specificity pocket (Lys802 for PI3K-γ versus Thr750 for PI3K-δ) as does the aminopyrazolopyrimidine at the hinge binding region^{21,22} (see Supporting Information, Figure S1).

Various substituted alkyne analogs at the 8-position that extended further into the non-conserved pocket were then prepared and evaluated for PI3K- γ potency and selectivity (Table 3). Additionally, as alkyl substituted alkyne analogs **16** and **17** suffered from poor in vitro metabolic stability, we also screened these substituted alkyne analogs for improved stability in mouse hepatocytes. Interestingly, hydrophobic substituents that extended deeper into the pocket (e.g. phenyl analog **18** vs. methyl

Table 2. Structure activity relationship of C8 substitution.



ACS Medicinal Chemistry Letters



^{*a*} Data reported as the average of at least two runs. ^{*b*}n=1. See Supporting Information for description of assay conditions and SEM values.

analog 17) were found to have eroded PI3K-γ potency and selectivity. Less hydrophobic groups (e.g. pyridyl analogs 19-21) maintained good selectivity for PI3K-γ and were found to be more metabolically stable than methyl alkyne analog 17. With this in mind, we synthesized and evaluated various analogs with smaller heterocycles on the end of the 8-alkyne (e.g. 22-26). Amongst various 5-membered ring heterocycle analogs, *N*-methylpyrazole analog 26 proved to have excellent potency and selectivity for PI3K-γ over PI3K-δ in enzymatic assays and excellent in vitro metabolic stability. Having achieved this remarkable selectivity in biochemical assays and in vitro and in vivo settings.

Compound 26 was evaluated for activity across all Class I PI3K isoforms. The binding affinity of compound 26 for PI3K- γ was determined by measuring the individual rates constants (k_{off} and k_{on}) and for PI3K- α , β and δ using equilibrium fluorescent titration. Compound 26 was found to be a remarkably tight binder to PI3K- γ with a K_d of 290 pM and >58-fold weaker affinity for other Class I PI3K isoforms (Table 4). Additionally, compound 26 did not significantly inhibit a panel of 468 mutant and non-mutant protein and lipid kinases (including Class II PI3K isoforms) at 1 µM (Tables S1 and S4). In PI3K- α , - β , - γ , and - δ dependent cellular phospho-AKT assays, compound 26 demonstrated excellent PI3K- γ potency $(IC_{50} = 1.2 \text{ nM})$ and selectivity against other Class I PI3K isoforms (>146-fold). Furthermore, compound 26 dose dependently inhibited PI3K-y dependent bone marrow-derived macrophage (BMDM) migration in vitro (Figure 1).²⁸

Table 3. Structure-activity relationship of C-8 alkynyl substitution.



Cound	D	Biochem (nl	nical IC ₅₀ $(M)^a$	Mouse Hepatocyte	
Chipa	K	PI3K-γ	PI3K-δ	Stability t _{1/2} (min)	
18		280	140	ND	
19	N N	110	>6000	102	
20	N N	350	>9000	51	
21	N	170	>9600	72	
22	N S	110	>8000	43	
23	S N	75	4800	270	
24	N S	360	>10000	27	
25	N N	100	>9000	251	
26	N-N	16	>8400	>360	

^{*a*} Data reported as the average of at least two runs. See Supporting Information for description of assay conditions and SEM values.

Table 4.	Class I	PI3K	selectivity	profile	for	compound	26	in
biochemi	cal and	cellula	r assays.					

A	PI3K Isoform						
Assay	α	β	γ	δ			
$K_{d}(nM)^{a}$	17	82	0.29	23			
Biochemical IC ₅₀ (nM)	3200	3500	16	>8400			

$(nM)^b$ 250 240 1.2 18	$\begin{array}{c} \text{Cellular IC}_{50} \\ (\text{nM})^b \end{array}$	250	240	1.2	180
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^{*a*} Binding affinities (K_d) of **26** were determined by methods previously reported.²⁴ See Supporting Information for full details. ^{*b*} Cellular IC₅₀s for Class I PI3K-α, PI3K-β, PI3K-γ, PI3K-δ were determined in SKOV-3, 786-O, RAW 264.7, and RAJI cells, respectively, by monitoring inhibition of pAKT S473 by ELISA.

Compound **26** was also found to be selective against a panel of 80 GPCRs, ion channels, and transporters at 10 μ M (see Supporting Information, Table S5 for full selectivity details). Additional in vitro safety assessment of compound **26** demonstrated that it was negative in Ames mutagenicity assays and had an IC₅₀ greater than 10 μ M in a hERG binding assay (data not shown).



Figure 1. Effect of compound **26** on migration of bone marrow derived macrophages (BMDM) in vitro. BMDMs were stimulated to migrate toward 100 μ g/mL CXCL12 (SDF-1 α) for 3 hours through a 5 micron Boyden Chamber with or without DMSO or a dose response of compound **26**. Migrated BMDMs were counted and compared to control.

In vitro absorption, distribution, metabolism and excretion (ADME) properties and pharmacokinetic parameters of compound **26** were also determined (summarized in Table 5). In vitro, **26** showed moderate to high cell permeability across Caco-2 cell monolayers, was slowly metabolized in cultured hepatocytes ($t_{1/2} > 360$ minutes), and demonstrated IC₅₀s greater than 20 μ M for the CYP isoforms tested (1A2, 2B6, 2C8, 2C9, 2C19, 2D6, 3A4). In vivo (mice, rats, dog, and monkeys), compound **26** had excellent oral bioavailability, low clearance, and distributed into tissues with a mean volume of distribution of 1.2 L/kg (Table 5). Overall, compound **26** had a favorable pharmacokinetic profile to allow potent and selective inhibition of PI3K- γ in vivo.

Based on the pharmacokinetic properties in mice and pharmacological in vitro profile, compound **26** was well-suited to investigate the impact of potent and selective PI3K- γ inhibition in vivo. IL-8 stimulated neutrophil migration into air pouches in mice has previously been shown to be dependent on PI3K- γ . Thus, to demonstrate PI3K- γ dependent activity of compound **26** in vivo, we evaluated the effect of orally administered compound **26** on IL-8 stimulated neutrophil influx into the air pouches on mice.^{7,8} Compound **26** significantly reduced neutrophil migration in a dose dependent manner in this model when administered orally at all of the tested doses (Figure 2a). The degree of inhibition observed directly correlated with plasma concentrations of compound **26** in these mice (Figure 2b), clearly demonstrated that orally administered compound **26** can inhibit PI3K- γ function in vivo. In addition, compound **26** (IPI-549) has been shown to inhibit tumor growth in murine syngeneic models through alteration of immune cells in the tumor microenvironment.²⁸

On the basis of its remarkable PI3K- γ potency and selectivity, favorable in vitro safety and pharmacokinetic profiles, and ability to inhibit PI3K- γ in vivo, compound **26** (IPI-549) was chosen as a development candidate. IPI-549 is currently in Phase 1 clinical evaluation in subjects with advanced solid tumors.²⁹

Table 5. In vitro ADME and pharmacokinetics for compound 26.

In vitro ADME							
Caco-2 ($P_{app} 10^{-6}$ cm/sec; A to B @ 10 μ M) 13							
$ \begin{array}{llllllllllllllllllllllllllllllllllll$							
Metabolic stability $(t_{1/2}; \min)^a$ > 360							
CYP3A4 inhibition (IC ₅₀ ; μ M) > 20							
Pharmacokinetics							
Mouse Rat Dog Monkey							
$C_{max}(\mu M)^b$	3.6	2.0	1.9	0.56			
$AUC_{0-\infty}$ $(ng*h/mL)^b$	20568	8049	1066	4030			
$t_{1/2}(h)^{c}$	3.2	4.4	6.7	4.3			
CL (mL/min/kg) ^c	3.6	4.4	2.8	4.3			
$V_{ss} (L/kg)^c$	0.8	1.2	1.3	1.3			

^a Study conducted using cultured human hepatocytes.

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^b Determined from oral dosing to male animals; PO dose levels: 5 mg/kg mouse and rat, 2.5 mg/kg monkey and dog. Formulation; 0.5% Carboxymethylcellulose, 0.05% Tween80 ^c Determined from intravenous dosing to male animals: IV dose levels: 1 mg/kg mouse and rat, 0.5 mg/kg monkey and dog. Formulation; 5% NMP, 40% PEG400, 55% PBS





Figure 2. (a) Effect of compound **26** on neutrophil migration in the mouse air pouch model. Mice were dosed orally one hour before IL-8 was applied to the air pouch. After 4 hours the air pouch was lavaged and the neutrophils counted using a CELL-DYN hematology analyzer. *indicates p values <0.05 compared to vehicle. Error bars indicate the standard error of the mean of 10 mice per group. (b) Plasma concentration levels of compound **26** at 1 h and 5 h post administration with overlaying cellular IC₅₀ (PI3K- γ and PI3K- δ) and IC₂₅ (PI3K- γ) values.

ASSOCIATED CONTENT

Supporting Information

The Supporting Information is available free of charge on the ACS Publications website at DOI: XXX

Synthetic procedures and analytical data for **1-26**; PI3K biochemical and cellular assay conditions; PI3K Class I & II selectivity data for **1** and **2**; full selectivity data for **26**; general animal study protocols (PDF)

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Notes

The authors declare no competing financial interest.

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ABBREVIATIONS

ADME, absorption, distribution, metabolism, and excretion; AKT, protein kinase B; ADP, adenosine diphosphate; ATP, adenosine triphosphate; BMDMs, bone marrow-derived macrophages; CXCL12, chemokine C-X-C motif 12; CYP, cytochrome P450; DMSO, dimethysulfoxide; ELISA, enzyme-linked immunosorbent assay; GPCR, G-protein-coupled receptor; IL-8, interleukin 8; NMP, *N*-methyl-2-pyrrolidone; PBS, phosphate-buffered saline; PI3K, phosphoinositide 3-kinase; PPB, plasma protein binding; PEG, polyethylene glycol; SDF-1, stromal cell-derived factor 1

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Page 7 of 7

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26 (IPI-549)