

8. Publications and Posters

List of Publications

1. Rajesh Bahekar, **Bhushan Dave**, Shubhangi Soman, Dipam Patel, Rajendra Chopade, Radhika Funde, Jeevan Kumar, S. Sachchidanand, Poonam Giri, Abhijit Chatterjee, Jogeswar Mahapatra, Purvi Vyas, Krishnarup Ghoshdastidar, Debduitta Bandyopadhyay, Ranjit C. Desai. Discovery of 1,3-dihydro-2H-imidazo[4,5-c]quinolin-2-ones based novel, potent and PI3K δ selective inhibitors. *Bioorganic & Medicinal Chemistry Letters* 29 (2019) 1313–1319.

List of Posters

1. **Bhushan Dave**, Rajendra Chopade, Dipam Patel, Jignesh Pethani, Hardik shah, Amitgiri Goswami, Abhijit Chatterjee, Purvi Vyas, Debduitta Bandyopadhyay, Jeevan Kumar, Mukul Jain, Rajesh Bahekar, Ranjit Desai, Shubhangi Soman. Design and Synthesis of Benzofuran based pyrazolo[3,4-*d*]pyrimidinamine derivatives as potent and selective PI3K δ Inhibitor. Poster presented in **8th RBF international Symposium, Advances in New Drug Discovery Technologies and Translational Research at Zydus Research Centre, Ahmedabad on Feb 2-4, 2017.**

2. **Bhushan Dave**, Rajendra Chopade, Dipam Patel, Jignesh Pethani, Hardik shah, Amitgiri Goswami, Abhijit Chatterjee, Purvi Vyas, Debduitta Bandyopadhyay, Jeevan Kumar, Sachchidanand S, , Mukul Jain, Rajesh Bahekar, Rajiv Sharma, Ranjit Desai, Shubhangi Soman. Design and Synthesis of 2,4-Disubstitued quinoline based pyrazolo [3,4-*d*] pyrimidinamine derivatives as potent and selective PI3K δ inhibitors. Poster presented in **9th RBF international Symposium, Advances in New Drug Discovery and Development at Zydus Corporate Park, Ahmedabad on Feb 6-8, 2020.**

3. Bhushan Dave, Rajendra Chopade, Dipam Patel, Amitgiri Goswami, Abhijit Chatterjee, Purvi Vyas, Debducta Bandyopadhyay, Sachchidanand S, Jeevan Kumar, Mukul Jain, Rajesh Bahekar, Rajiv Sharma, Ranjit Desai, Shubhangi Soman. Design, Synthesis and Pharmacological evolution of Imidazo-Quinolinyll derivatives as PI3Kdelta-Selective Inhibitors. Poster presented in **Advances in Chemistry of Bioactive Molecules (ACBAM-2020) at M.S. University, Baroda on Jan 17-18, 2020**. Above poster presentation was awarded 1st prize.



Contents lists available at ScienceDirect

Bioorganic & Medicinal Chemistry Letters

journal homepage: www.elsevier.com/locate/bmclDiscovery of 1,3-dihydro-2*H*-imidazo[4,5-*c*]quinolin-2-ones based novel, potent and PI3K δ selective inhibitors[☆]

Rajesh Bahekar^{a,*}, Bhushan Dave^{a,b}, Shubhangi Soman^b, Dipam Patel^a, Rajendra Chopade^a, Radhika Funde^a, Jeevan Kumar^c, S. Sachchidanand^e, Poonam Giri^d, Abhijit Chatterjee^d, Jogeswar Mahapatra^d, Purvi Vyas^e, Krishnarup Ghoshdastidar^e, Debdutta Bandyopadhyay^e, Ranjit C. Desai^a

^a Department of Medicinal Chemistry, Zydus Research Centre, Sarkhej-Bavla, N.H. 8A Moraiya, Ahmedabad 382210, India

^b Department of Chemistry, Faculty of Science, M.S. University of Baroda, Vadodra 390002, India

^c Department of Bioinformatics, Zydus Research Centre, Sarkhej-Bavla, N.H. 8A Moraiya, Ahmedabad 382210, India

^d Department of Pharmacology, Zydus Research Centre, Sarkhej-Bavla, N.H. 8A Moraiya, Ahmedabad 382210, India

^e Department of Cell Biology, Zydus Research Centre, Sarkhej-Bavla, N.H. 8A Moraiya, Ahmedabad 382210, India

ARTICLE INFO

Keywords:
PI3K inhibitors
Inflammatory and autoimmune diseases
Rheumatoid arthritis
Imidazo-quinolinones
Design
Synthesis and biological study

ABSTRACT

PI3K δ is implicated in various inflammatory and autoimmune diseases. For the effective treatment of chronic immunological disorders such as rheumatoid arthritis, it is essential to develop isoform selective PI3K δ inhibitors. Structure guided optimization of an imidazo-quinolinones based pan-PI3K/*m*-TOR inhibitor (Dactolisib) led to the discovery of a potent and orally bioavailable PI3K δ isoform selective inhibitor (**10h**), with an improved efficacy in the animal models.

Rheumatoid arthritis (RA) is a chronic inflammatory disease characterized by synovitis and joint destruction. Treatment with biologic agents such as tumor necrosis factor (TNF) inhibitors has improved outcomes, but in general, biologics are expensive, injectable and many patients have inadequate responses. Thus, orally bioavailable small molecule inhibitors that target signal transduction and regulate innate and adaptive immune responses in RA have emerged as potential alternatives to expensive biologics.¹

Phosphoinositide-3-kinases (PI3Ks) constitute central signaling hub that mediates diverse and crucial cell functions, including cell growth, proliferation, differentiation and survival.^{2,3} PI3Ks have been classified into three classes (I, II and III) based on substrate specificity, sequence homology and regulatory subunits. The class I PI3Ks consists of four kinases (PI3K- α , β , δ and γ) and further grouped into two sub-classes: class IA and class IB. The class IA comprises three closely related kinases, PI3K- α , β and δ , while the class IB contains only one member PI3K- γ .⁴ The PI3K α and β are expressed in a wide variety of tissues and organs. PI3K γ is found mainly in leukocytes, while expression of PI3K δ is restricted to spleen, thymus, hematopoietic cells and peripheral blood leukocytes.⁵ PI3K γ and PI3K δ are mainly expressed in rheumatoid

arthritis (RA) synovium and regulate innate and adaptive immune responses.⁶

Inhibition of PI3Ks is considered as one of the most interesting targets. Earlier attempts were mainly focused on developing the broad-spectrum (pan) inhibitors of the PI3K (α , β , γ and δ) isoforms, as potential oncology therapeutics.^{7–9} Knowing the potential side effects associated with PI3K α and β isoforms inhibition (due to universal expression), recently, efforts are directed towards the development of PI3K δ selective inhibitors, for the effective treatment of hematological malignancy and inflammatory disorders.^{10–12}

Over the past decades, several structurally diverse PI3K inhibitors were identified containing quinolines and imidazoquinolinones as promising scaffold (Fig. 1). The Omipalisib^{13,14} and Dactolisib¹⁴, discovered by GlaxoSmithKline and Novartis respectively, as dual PI3K and *m*TOR inhibitors are under clinical trials. PI3K δ selective inhibitors, Idelalisib¹⁵ (ZYDELIG[®], Gilead Sciences, in 2014) is available for the treatment of hematologic malignancies. Recently, US-FDA approved Duvelisib^{16,17} (COPIKTRA[®], Verastem, Inc) for the treatment of chronic lymphocytic leukemia (CLL) or small lymphocytic lymphoma (SLL). There is still requirement to develop safe and highly potent PI3K δ

[☆] ZRC communication No: 587 (Part of PhD thesis work of Mr. B. Dave).

* Corresponding author.

E-mail address: rajeshbahekar@zyduscadila.com (R. Bahekar).

<https://doi.org/10.1016/j.bmcl.2019.04.007>

Received 24 December 2018; Received in revised form 19 March 2019; Accepted 3 April 2019

Available online 04 April 2019

0960-894X/ © 2019 Elsevier Ltd. All rights reserved.

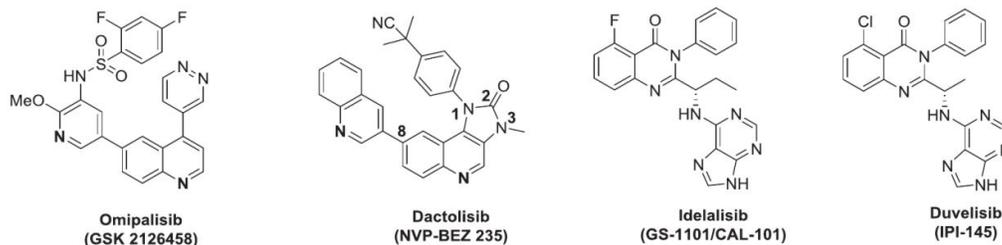


Fig. 1. The structures of quinoline and imidazoquinolinone-based PI3K inhibitors.

selective inhibitors, for the effective management of chronic inflammatory and autoimmune disorders, such as rheumatoid arthritis (RA) and hematological malignancy.

We aim to discover novel, potent and orally bioavailable PI3K δ selective inhibitors, mainly by favoring the suitable accommodation of designed molecules, in the specificity pocket, to achieve PI3K δ selectivity.¹⁸ Considering imidazoquinolinone moiety as a starting point, appropriate structural modifications were carried out in the Dactolisib (pan-PI3K/mTOR inhibitor), to improve PI3K δ selectivity. Two set of compounds were designed (as listed in Tables 1 and 2). Initial modifications on imidazole ring (of Dactolisib), involves positional changes of methyl (N1 to N3) and phenyl (N3 to N1) groups and introduction of a carbon spacer (phenyl to benzyl) at N3 position. Further modifications were carried out at the *p*-position of benzyl ring (Set-1, Table 1) to obtain the single digit nM potency (9c). Compound 9c was found to be potent but showed moderate isoform selectivity (Table 3). In the second set (Set-2, Table 2), changes were carried out on the 8th position of 9c to improve isoform selectivity and in vivo efficacy.

Herein, we report, design, synthesis and biological evaluation of novel 1,3-dihydro-2*H*-imidazo[4,5-*c*]quinolin-2-one based PI3K δ selective inhibitors. Test compounds were screened in vitro for PI3K δ inhibitory activity and most potent compounds from each set were

tested for in vitro PI3K isoform selectivity (α , β & γ) and mTOR inhibitory activity.¹⁹ Based on the in vitro results, highly potent and selective compound 10h was selected for in vivo PK and PD (anticancer and anti-inflammatory activities) studies.

Synthesis of 1,3-dihydro-2*H*-imidazo[4,5-*c*]quinolin-2-one derivatives (9a-g and 10a-m) was carried out as depicted in Scheme 1, following the modified literature procedure.²⁰ Treatment of bromo anthranilic acid (1) with nitro methane gives nitro vinyl anthranilic acid (2), which was cyclized using potassium acetate in acetic anhydride to get the bromo nitro quinolinol (3). Compound 3 was converted to reactive chloro derivate (4) using POCl₃, followed by nucleophilic substitution with methylamine to get the compound 5. Nitro group of 5 was reduced, using SnCl₂ to get the compound 6, which was cyclized using diphosgene in the presence of base, to obtain the imidazoquinolinone (7). Alkylation of 7 using strong base and aryl halides furnished compounds 8a-g, which were converted to 9a-g and 10a-m, using PdCl₂(PPh₃)₂, potassium bicarbonate and aryl or heteroaryl boronic acids.

Overall, 20 compounds (9a-g and 10a-m) were prepared in good yield (60 to 80%), under the mild reaction condition. Spectral data of compounds were found to be in conformity with the structures assigned, which ensure the formation of the compounds 9a-g and 10a-m (see supporting information for analytical and spectral data).

For in vitro PI3K δ inhibitory activities, Idelalisib and Dactolisib were used as a positive control. In Set-1 (Table 1), 4-substituted-(benzyl)-8-quinolinyl-imidazo[4,5-*c*]quinolinone (9a-g) analogues displayed varying degree of PI3K δ inhibitory activities at 100 nM concentration. Compound 9a (2-methyl-propanenitrile substitution on a benzyl ring) showed moderate PI3K δ inhibitory activity (56% PI3K δ inhibition at 100 nM), while replacement of the nitrile group (Compound 9b: 73% inhibition) exhibited enhanced PI3K δ inhibitory activity (IC₅₀: 28.3 nM). Replacement of isopropyl (9b) with methoxy (9c, IC₅₀: 9.5 nM) and methyl (9d, IC₅₀: 18.4 nM) demonstrated higher PI3K δ inhibitory activity, whereas replacement with an electron withdrawing NO₂ (9e, 62% inhibition), electronegative F (9f: 51% inhibition) and the unsubstituted compound 9g (22% inhibition) displayed weak PI3K δ activity.

In Set-1, compound 9c (methoxy) showed most potent PI3K δ inhibitory activity, which was found to be similar to Dactolisib (IC₅₀: 8 nM; Table 1) and 4.5 fold less potent compared to Idelalisib (IC₅₀: 2.1 nM). Thus, in the Set-1, positional changes of methyl (N1 to N3) and phenyl (N3 to N1) groups and introduction of a carbon spacer (phenyl to benzyl) at N3 position, followed by substitution on *p*-position of benzyl ring led to the single digit nM potent compound (9c) with moderate isoform selectivity (Table 3).

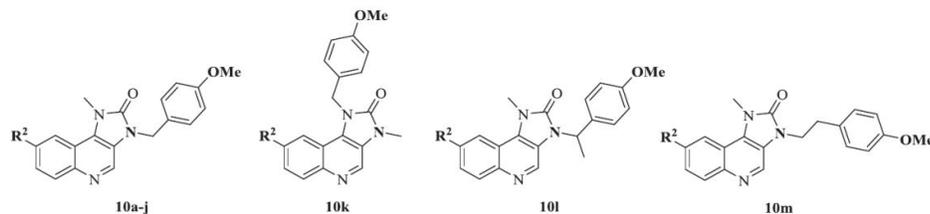
To improve PI3K δ isoform selectivity, modifications were carried out in 9c as a primary lead. In the second set (Table 2), we found that the PI3K δ inhibitory activity was retained by doing modifications at 8th position of quinoline in 9c. As listed in Table 2 (Compounds 10a-m), replacement of quinoline moiety in 9c with benzothiazole (10a and 10b (acylated 10a)) showed 50% PI3K δ inhibition at 100 nM. Phenyl

Table 1
PI3K δ inhibitory activity of 4-substituted-(benzyl)-8-quinolinyl-imidazo[4,5-*c*]quinolinone (9a-g).

| Comp. | R ¹ | PI3K δ inhibition (%) ^{a,b} | PI3K δ IC ₅₀ (nM) ^c |
|------------|------------------|---|--|
| 9a | | 56 | ND |
| 9b | | 73 | 28.3 |
| 9c | | 98 | 9.5 |
| 9d | -CH ₃ | 80 | 18.4 |
| 9e | -NO ₂ | 62 | ND |
| 9f | -F | 51 | ND |
| 9g | -H | 22 | ND |
| Dactolisib | - | 92 | 8 |
| Idelalisib | - | 96 | 2.1 |

^aAll the data are shown as the mean for at least two experiments. ^bPI3K δ inhibition at the concentration of 100 nM using PI3K Kinase Activity/Inhibitor assay Kit from Millipore. ^cThe IC₅₀ values for PI3K δ inhibition. ND: not detected.

Table 2

Influence of modification of C8 position of Quinoline moiety on PI3K δ isoform selectivity and mTOR selectivity.

| Comp. | R ² | PI3K δ inhibition (%) ^{a,b} | PI3K δ IC ₅₀ (nM) ^c | Comp. | R ² | PI3K δ inhibition (%) ^{a,b} | PI3K δ IC ₅₀ (nM) ^c |
|-------|----------------|---|--|------------|----------------|---|--|
| 10a | | 52 | ND | 10h | | 99 | 1.9 |
| 10b | | 58 | ND | 10i | | 91 | 6.2 |
| 10c | | 72 | 29.4 | 10j | | 93 | 5.9 |
| 10d | | 81 | 20 | 10k | | 98 | 4.1 |
| 10e | | 87 | 11.4 | 10l | | 60 | ND |
| 10f | | 88 | 11.1 | 10m | | 56 | ND |
| 10g | | 62 | ND | Idelalisib | | 96 | 2.1 |

^aAll the data are shown as the mean for at least two experiments. ^bPI3K δ inhibition at the concentration of 100 nM using PI3K Kinase Activity/Inhibitor assay Kit from Millipore. ^cThe IC₅₀ values for PI3K δ inhibition. ND: not detected.

Table 3

Isoform selectivity of compounds against PI3K (α , β , γ , and δ) and mTOR activities.

| Comp. | Biochemical IC ₅₀ (nM) ^a | | | | |
|------------|--|---------------------------|----------------------------|----------------------------|----------------------------|
| | PI3K α ^b | PI3K β ^b | PI3K γ ^b | PI3K δ ^b | mTOR ^b (p70S6K) |
| 9c | 421 | 342 | 38 | 9.5 | 676 |
| 10h | 891 | 589 | 112 | 1.9 | > 1000 |
| 10k | 289 | 241 | 42 | 4.1 | 580 |
| Dactolisib | 5 | 79 | 6 | 8 | 14 |
| Idelalisib | 831 | 571 | 92 | 2.1 | > 1000 |

^aThe IC₅₀ values are shown as the mean for at least two experiments. ^bPI3K inhibitory activity assay Kit (Millipore) was used to screen the test compounds.

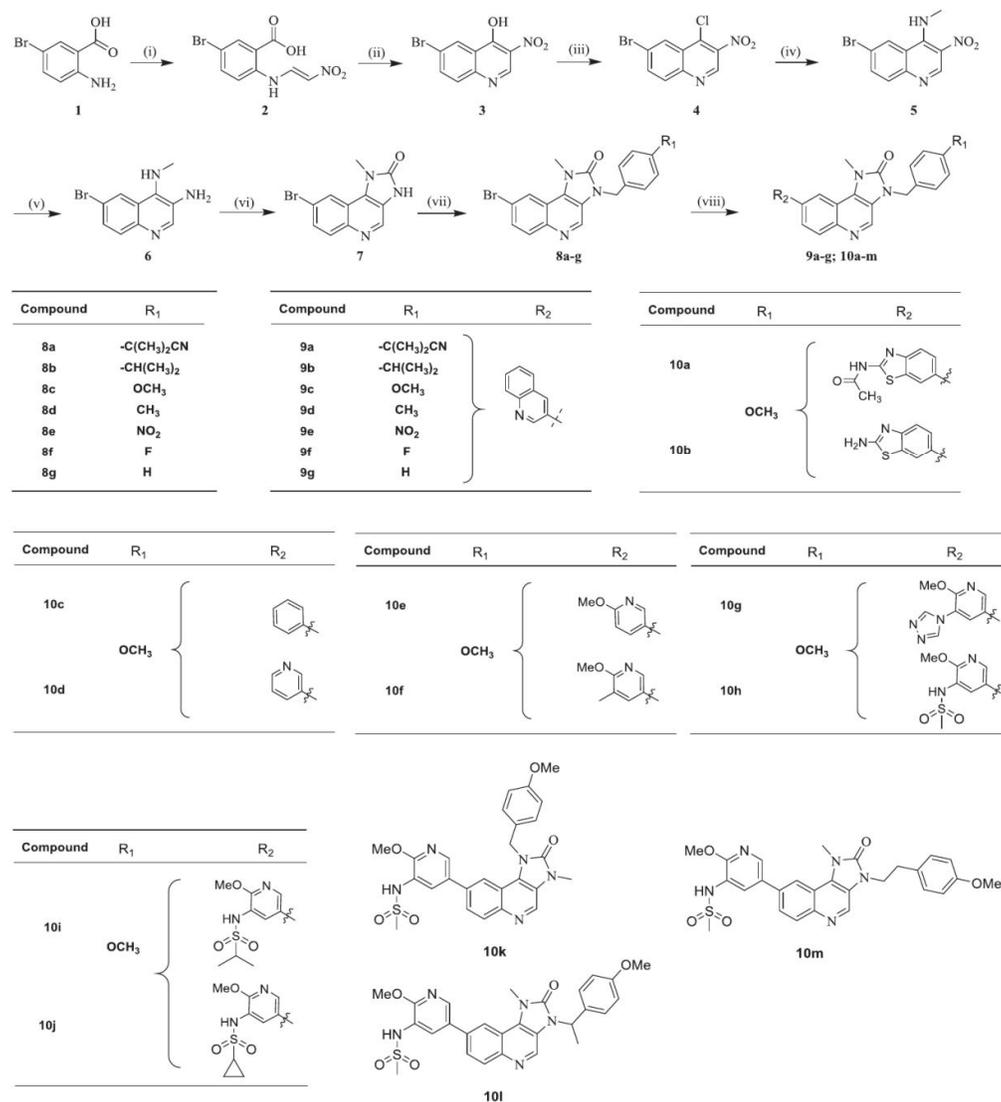
derivative (**10c**) was found to be less potent (PI3K δ IC₅₀: 29.4 nM) compared to **9c**, while Pyridyl derivatives (**10d**) showed some improvement in the potency (IC₅₀: 20 nM). Introduction of methoxy (**10e**) and 3-methyl-2-methoxy substituents (**10f**) were found to be equipotent (IC₅₀: ~11 nM). Triazole motif (**10g**) showed less activity compared to **9c**, while *m*-methanesulfonamide derivative (**10h**) was found to be the

most potent (IC₅₀: 1.9 nM) among all. Replacement of methyl group in **10h**, with isopropyl (**10i**) and cyclopropyl (**10j**) showed three fold less activity compared to **10h**.

Compound **10k**, a regioisomer of **10h**, wherein benzyl and methyl groups are interchanged on imidazole ring showed two fold less active compared to **10h**. Similarly, racemic compound **10l**, and **10m** (one carbon homologs at N3 with respect to **10h**) were found to be less active.

Most potent compounds (**9c**, **10h** and **10k**) were evaluated for their selectivity against PI3K isoforms (α , β and γ) and mTOR. As shown in the Table 3, initial hits (**9c** and **10k**) showed moderate selectivity against PI3K isoforms and mTOR over PI3K δ . Compound **10h** (IC₅₀: 1.9 nM) demonstrated 469, 310, and 59-fold selectivity over PI3K α , β and γ respectively. Moreover, it was noted that selectivity of **10h** against all the three isoforms was higher than standard compounds. In general, it was observed that the potency and selectivity of imidazoquinolinone-based PI3K δ inhibitors can be modulated using suitable substituents at R¹ and R² positions.

In vitro kinase profiling study of **10h** was carried out @ 1 μ M concentration, against 140 kinases and % inhibition was found to be < 20% at 1 μ M concentration. Compound **10h** was tested for its anti-



Scheme 1. Synthesis of compounds **9a-g** and **10a-l**. Reagents and conditions: i) conc. HCl, water, 26 °C, 6h, then nitro methane, NaOH, water, Conc. HCl, 26 °C, 16 h, 95%; ii) KOAc, acetic anhydride, 120–125 °C, 4h, 60%; iii) POCl₃, 120 °C, 4 h, 80%; iv) Me-NH₂, TEA, DCM, 26 °C, 12 h, 94%; v) SnCl₂·2(H₂O)₂, EtOAc, 26 °C, 60%; vi) Diphosgene, TEA, DCM, 0–26 °C, 65%; vii) R₁-Ar-CH₂-X (Cl or Br), NaH, THF, 0–26 °C, 65–75%; viii) R₂-B(OH)₂, PdCl₂(PPh₃)₂, KHCO₃, DMF, H₂O, 90–95 °C, 1.5 h, 60–80%.

proliferative activities against TMD-8 cell lines²¹ with Idelalisib as a reference compound. In anti-proliferative in vitro assay, **10h** and Idelalisib exhibited potent anti-proliferative activity with an IC₅₀ value of 340 and 795 nM respectively. Additional profiling studies of compound **10h** was carried out and it was found to be devoid of CYP²² (< 10% CYP inhibition at 10 μM concentration, for CYP1A2, CYP2C8, CYP2C9,

CYP2D6, CYP2C19 and CYP3A4) and hERG liabilities (IC₅₀: > 30 μM), while Idelalisib showed moderate CYP3A4 inhibition.²³

A comparative single dose (3 mg/kg, po and 1 mg/kg, iv) PK profile of compounds **9h**, **10h** and Dactolisib was evaluated in male C57BL/6J mice (n = 6) and the various PK parameters (T_{max}, C_{max}, t_{1/2}, Cl, AUC and %F) were recorded (Table 4). In PK study, **9c** showed moderate

Table 4
Pharmacokinetic study parameters^a of **9c**, **10h** and Dactolisib in C57 mice.

| Compd | Tmax (h) | Cmax (µg/ml) | t _{1/2} (h) | Cl (ml/min/kg), iv | AUC (0-α) h µg/ml | %F ^b |
|------------|----------|--------------|----------------------|--------------------|-------------------|-----------------|
| 9c | 0.38 | 127.61 | 2.31 | 22.9 | 329.08 | 14.77 |
| 10h | 0.25 | 1278.49 | 3.48 | 8.24 | 3831.48 | 68.91 |
| Dactolisib | 0.21 | 273.83 | 1.88 | 72.5 | 243.64 | 37.82 |

^aOral bioavailability (%F) was calculated wrt to iv AUC. Compounds **9c**, **10h** and Dactolisib administered at 1 mg/kg dose, iv AUC: 742.56, 1852.95 and 215.14 respectively.

^bIn male C57BL/6J mice (n = 6), compounds were administered orally (p.o) at 3 mg/kg dose and plasma concentration was analyzed by LC-MS, values indicate Mean ± SD.

AUC and clearance, which resulted into overall low bioavailability (~15%). Compound **10h** showed rapid Tmax, higher AUC (~5 fold, compared to standard), extended t_{1/2} (~3.5 hr) and good oral bioavailability (%F ~69 over standard, 38%). Compound **10h** showed extended t_{1/2} and higher AUC, which could be due to its low clearance compared to standard (8.24 vs 72.5 ml/min/kg, iv).

Considering low bioavailability of **9c**, in PD models, only **10h** was evaluated. Collagen Induced Arthritis (CIA) mice model was used to check anti-arthritis efficacy of test compounds.¹⁷ Arthritis was developed in male DBA1j mice, using collagen mixture and mice were recruited for the study once clinical signs were visible. Eight animals were assigned in each of the four groups [vehicle, positive control (Dactolisib, 60 mg/kg dose was selected considering low oral bioavailability, compared to **10h**) and two doses of test compound **10h** (10 and 30 mg/

kg) was selected considering 89% plasma protein binding and t_{1/2} of ~3.5 h]. Treatment was continued for four weeks and percentage inhibition in clinical score was recorded.

As shown in the Fig. 2a, standard and **10h** showed good reduction in the arthritic score, compared to vehicle control (untreated group). Two fold higher dose of a standard compound was used, considering two fold difference in the mice oral bioavailability. At 30 mg/kg dose, compound **10h** showed superior activity compared to standard compound (dose 60 mg/kg). Body weights of the animals were also recorded 3 times a week as a measure of treatment related side effect and **10h** showed no significant reduction in the body weight, even at 30 mg/kg dose, while Dactolisib exhibited reduction in body weight.

Additionally, in vivo anti-tumor activity of **10h** was checked in male SCID mice xenograft model (inoculated with TMD-8 cells). Inhibition of tumor volume compared to vehicle control (untreated group) was considered as efficacy end point. As shown in Fig. 2b, three doses (3, 10 and 30 mg/kg/day, orally) of **10h** were administered and it showed dose dependent reduction in the tumor volume. At 30 mg/kg dose, **10h** showed complete inhibition of tumor volume compared to vehicle control. Body weights of the animals were also recorded and **10h** showed no significant reduction in the body weight, even at 30 mg/kg dose. Thus improved PK of **10h** justifies its potent in vivo activity in both the animal models.

The structure of mPI3Kδ co-crystallised with Idelalisib (PDB ID: 4XE0)¹⁸ was prepared using protein preparation wizard of Schrodinger Suite 2018 at pH 7.4, which was used for generating the docking grid. The Glide docking was performed using Glide SP with default parameters. Ligand molecules (Idelalisib, Dactolisib, **9c**, **10h** and **10k**) were prepared using LigPrep module of Schrodinger Suite 2018.²⁴

The results of docking has been summarized in Fig. 3 where it is clear from the docking poses of the docked ligands that specificity pocket was fully occupied by **10h** and **10k**, however in case of Dactolisib and **9c**, the specificity pocket was not occupied and this explains why **10h** and **10k** were more selective towards PI3Kδ.

In conclusion, we have synthesized and evaluated two sets of novel series of 1,3-dihydro-2H-imidazo[4,5-c]quinolin-2-one derivatives as selective PI3Kδ inhibitors. In first set, appropriate modifications were carried out in the imidazoquinoline ring, which led to an identification of a single digit nM potent PI3Kδ inhibitor (**9c**), with moderate isoform selectivity. In set 2, further structure-activity relationship (SAR) studies on the 8th position of **9c** resulted in to the discovery of N-(2-methoxy-5-(3-(4-methoxybenzyl)-1-methyl-2-oxo-2,3-dihydro-1H-imidazo[4,5-c]quinolin-8-yl)pyridin-3-yl)methanesulfonamide (**10h**) that showed improved isoform selectivity, PK profile and good efficacy in a CIA and xenograft animal models. The results of docking study showed that specificity pocket was fully occupied by **10h**, which explains its more selective towards PI3Kδ. Overall pre-clinical data suggest that the development of a potent and selective PI3Kδ inhibitor could be viable therapeutic option for the effective management of rheumatoid arthritis and hematological malignancy.

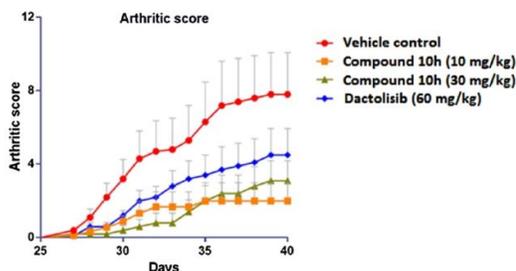


Fig. 2a. Effect of Compound **10h** and Dactolisib in CIA mice model.

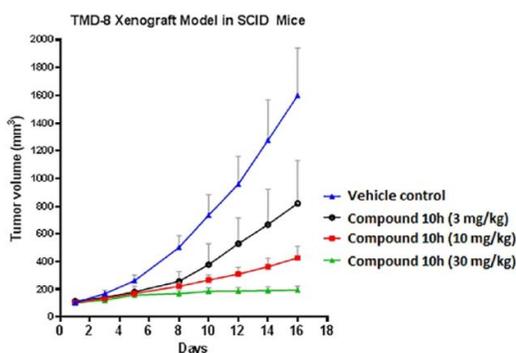


Fig. 2b. In vivo anti-tumor activity of Compound **10h** in SCID mice xenograft model.

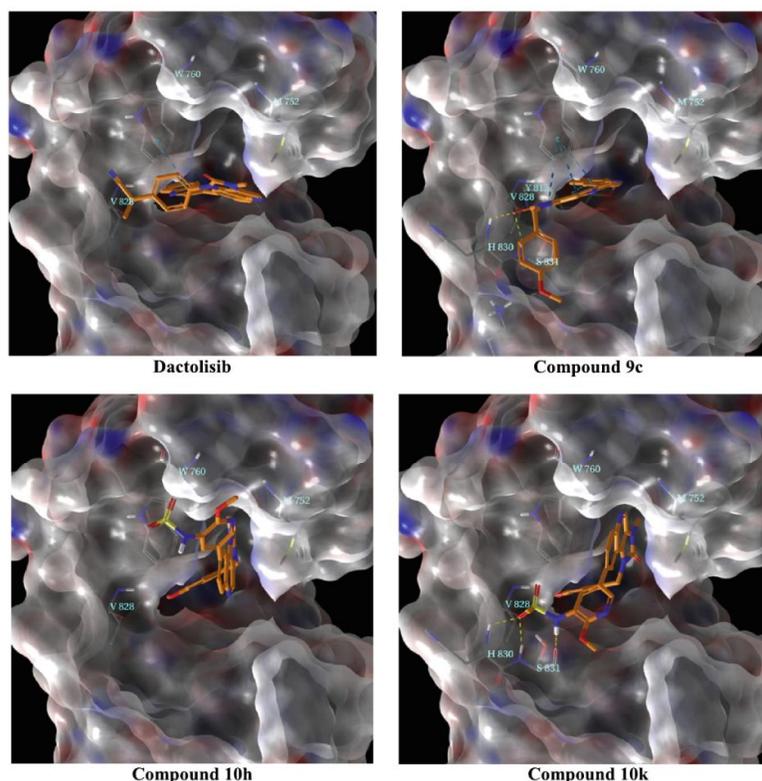


Fig. 3. The Glide docking studies of Compounds 9c, 10h, 10k and Dactolisib into the site of PI3K δ (PDB ID: 4XE0). Compounds are shown as sticks. Hydrogen bonds are shown as yellow dashed lines.

Acknowledgments

Authors thank the management of Zydus Research Centre, Cadila Healthcare Ltd. for support and encouragement and the analytical department for providing analytical data.

Notes

The authors declare no competing financial interest.

Appendix A. Supplementary data

Supplementary data to this article can be found online at <https://doi.org/10.1016/j.bmcl.2019.04.007>.

References

- David LB, Hae RK, Katharyn T, Beatrix B, Gary SF. Novel phosphoinositide 3-kinase δ , γ inhibitor: potent anti-inflammatory effects and joint protection in models of rheumatoid arthritis. *J Pharmacol Exp Ther.* 2014;348:271–280.
- Cantley LC. The phosphoinositide 3-kinase pathway. *Science.* 2002;296:1655–1657.
- Hawkins PT, Stephens LR. PI3K signalling in inflammation. *Biochim Biophys Acta Mol Cell Biol Lipid.* 1851;2015:882–897.
- Geering B, Cutillas PR, Nock G, Gharbi SI, Vanhaesebroeck B. Class IA phosphoinositide 3-kinases are obligate p85–p110 heterodimers. *PNAS.* 2007;104:7809–7814.
- Knight Z, Gonzalez B, Feldman M, et al. A pharmacological map of the PI3-K family defines a role for p110 α in insulin signaling. *Cell.* 2006;125:733–747.
- Bartok B, Boyle DL, Liu Y, et al. PI3 kinase δ is a key regulator of synovioyte function in rheumatoid arthritis. *Am J Pathol.* 2012;180:1906–1916.
- Marone R, Cmiljanovic V, Giese B, Wymann MP. Targeting phosphoinositide 3-kinase: moving towards therapy. *Biophys Acta Proteins Proteomics.* 2008;1784:159–185.
- Maira SM, Voliva C, Garcia-Echeverria C. Class IA phosphatidylinositol 3-kinase: from their biologic implication in human cancers to drug discovery. *Expert Opin Ther Targets.* 2008;12:223–238.
- Foster JG, Blunt MD, Carter E, Ward SG. Inhibition of PI3K signaling spurs new therapeutic opportunities in inflammatory/autoimmune diseases and hematological malignancies. *Pharmacol Rev.* 2012;64:1027–1054.
- Hirsch E, Katanaev VL, Garlanda C, et al. Central role for G protein-coupled phosphoinositide 3-kinase gamma in inflammation. *Science.* 2000;287:1049–1053.
- Li Z, Jiang H, Xie W, Zhang Z, Smrcka AV, Wu D. Roles of PLC-beta2 and -beta3 and PI3Kgamma in chemoattractant – mediated signal transduction. *Science.* 2000;287:1046–1049.
- Patrucco E, Notte A, Barberis L, et al. PI3Kgamma modulates the cardiac response to chronic pressure overload by distinct kinase-dependent and -independent effects. *Cell.* 2004;118:375–387.
- Steven DK, Nicholas DA, Joelle LB, et al. Discovery of GSK2126458, a highly potent inhibitor of PI3K and the mammalian target of rapamycin. *ACS Med Chem Lett.* 2010;1:39–43.
- Hua-fu Z, Jing W, Wei S, et al. Recent advances in the use of PI3K inhibitors for glioblastoma multiforme: current preclinical and clinical development. *Mol Cancer.* 2017;16:100–115.
- John RS, David K, Armando GV, et al. Structural biochemical and biophysical characterization of idelalisib binding to phosphoinositide 3-kinase δ . *J Biol Chem.* 2015;290:8439–8446.
- Genevra P, Niamh VL, Rachel EP, et al. Targeting PI3K δ and PI3K γ signalling disrupts human AML survival and bone marrow stromal cell mediated protection. *Oncotarget.* 2016;7:39784–39795.
- David GW, Kerrie LF, Jonathan PD, et al. PI3K- δ and PI3K- γ inhibition by IPI-145 abrogates immune responses and suppresses activity in autoimmune and inflammatory disease models. *Chem Biol.* 2013;20:1364–1374.
- Somoza R, Koditek D, Villaseñor G, et al. Structural, biochemical, and biophysical characterization of idelalisib binding to phosphoinositide 3-kinase delta. *J Biol Chem.* 2015;290:8439–8446.

19. Christopher S, Kelley LR, Colleen WO, Esther PB. Inhibition of class IA PI3K enzymes in non-small cell lung cancer cells uncovers functional compensation among isoforms. *Cancer Biol Ther.* 2015;16:1341–1352.
20. Stowasser F, Banziger M, Garad S D. Salts and crystal forms of 2-methyl-2-[4-(3-methyl-2-oxo-8-quinolin-3-yl-2,3-dihydro-imidazo[4,5-c]quinolin-1-yl)-phenyl]-propanitrile, WO 2008064093.
21. Juliane P, Maurice S, Antje MW, et al. Simultaneous inhibition of PI3K δ and PI3K α induces ABC-DLBCL regression by blocking BCR-dependent and -independent activation of NF- κ B and AKT. *Cancer Cell.* 2017;31:64–78.
22. Kumar GN, Surapaneni S. Role of drug metabolism in drug discovery and development. *Med Res Rev.* 2001;21:397–411.
23. Jin F, Robeson M, Zhou H, et al. Clinical drug interaction profile of idelalisib in healthy subjects. *J Clin Pharmacol.* 2015;55:909–919.
24. Schrödinger Release, 2018-3: Glide, Schrödinger, LLC, New York, NY, 2018.

Design and Synthesis of Benzofuran based pyrazolo[3,4-d]pyrimidinamine derivatives as potent and selective PI3K δ inhibitors

Bhushan Dave^{1,2}, Rajendra Chopade³, Dipan Patel⁴, Jignesh Pethani⁵, Hardik Shah⁶, Amitgiri Goswami⁷, Abhijit Chatterjee⁸, Purvi Vyas⁹, Debudutta Bandyopadhyay¹⁰, Jeevan Kumar¹¹, Mukul Jain¹², Rajesh Babekar¹³, Ranjit Desai¹⁴ and Shubhangi Soman¹⁵

¹Department of Medicinal Chemistry, ²Department of Pharmacology, ³Department of Cell Biology, ⁴Zydus Research Centre, Sarkhej-Bavla N. H. No. RA, Morbiya, Ahmedabad 382210, Gujarat, India
⁵Department of Chemistry, Faculty of Science, M.S. University of Baroda, Vadodra 390002, India

Abstract
Phosphoinositide-3-kinases (PI3Ks) mediate diverse and crucial cell functions. The class IA PI3Ks comprises PI3K- α and β , while the class IB contains PI3K- δ . Knowing the potential side effects associated with PI3K- α and β isoforms inhibition (due to universal expression), recent efforts are focused towards the development of potent and PI3K- δ selective inhibitors for the treatment of hematologic malignancies and inflammatory diseases. In the present investigation, we report synthesis and *in vitro* screening of benzofuran based pyrazolo[3,4-d]pyrimidinamine derivatives (10a-j) as potent and selective PI3K δ inhibitors. Further test compounds were screened for mTOR selectivity and compounds 10a-c & 10h-j exhibited better PI3K δ selectivity over mTOR.

Introduction
PI3K constitute central signaling hub that mediates many diverse and crucial cell functions, including cell growth, cell proliferation, metabolism and survival. Isoforms, p110 α and p110 β are ubiquitously expressed, while p110 δ and p110 ϵ are expressed predominantly in hematopoietic cells, especially, PI3K δ is expressed in leukocytes and considered as useful target for the treatment of hematologic malignancies and inflammatory diseases. Several groups are working for identifying selective inhibitors of PI3K δ . Currently a number of selective PI3K δ inhibitors are discovered, which are in various stages of clinical development for diverse indications such as cancer, asthma and inflammation.

Mechanism of Action
PI3Ks are lipid kinases that catalyze the phosphorylation of 3'-hydroxyl group of phosphatidylinositol and phosphatidylinositides in response to extracellular stimuli. The 3-phosphorylated phospholipids (PPLs) generated by PI3K act as second messengers, recruiting kinases with lipid binding domains, such as Akt and PDK1. Binding of Akt to membrane PPLs causes the translocation of Akt to the plasma membrane and its activation.

Results and Discussion
Title compounds 10a-j were designed as a bioisosteric replacement of quinoline ring of INK-654 with benzofuran ring system (Figure 1). The synthesis of titled compounds 10a-j was carried out as depicted in Scheme 1, followed by *in vitro* testing in PI3K, mTOR assays of p110 δ and p110 ϵ . Among the compounds tested, 10a-c & 10h-j showed good *in vitro* PI3K δ inhibitory activity (IC_{50} inhibition at 10 nM). The test compounds were also screened for mTOR selectivity and compounds 10a-c & 10h-j exhibited better PI3K δ selectivity over mTOR. Additionally, molecular docking studies were carried out for 10j and the docking results of 10j correlates with its potent *in vitro* PI3K δ activity.



Figure 1: PI3K activation pathways

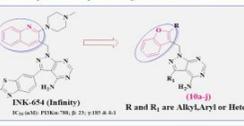
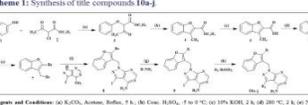


Figure 2: Bioisosteric replacement of quinoline ring of INK-654 with benzofuran ring system



Figure 3: Binding pose of 10j in PI3K δ specificity pocket

Experimental
Scheme 1: Synthesis of title compounds 10a-j



Reagents and Conditions: (a) K_2CO_3 , Acetone, Reflux, 1 h; (b) CH_2N_2 , CH_2Cl_2 , 0°C to RT, 1 h; (c) $NaHCO_3$, CH_2Cl_2 , 0°C to RT, 1 h; (d) $NaHCO_3$, CH_2Cl_2 , 0°C to RT, 1 h; (e) $NaHCO_3$, CH_2Cl_2 , 0°C to RT, 1 h; (f) $NaHCO_3$, CH_2Cl_2 , 0°C to RT, 1 h; (g) $NaHCO_3$, CH_2Cl_2 , 0°C to RT, 1 h; (h) $NaHCO_3$, CH_2Cl_2 , 0°C to RT, 1 h; (i) $NaHCO_3$, CH_2Cl_2 , 0°C to RT, 1 h; (j) $NaHCO_3$, CH_2Cl_2 , 0°C to RT, 1 h.

Chemistry: Overall, 10 compounds (10a-j) were prepared in good yield, under the mild reaction condition (Scheme 1) and were characterized with their physical, analytical and spectral data (1H NMR, ESI MS). The percentage yield in the final step was found to be in the range of 60-80%. The ESI MS and NMR spectral data of all the synthesized compounds were also found to be in conformity with the structures assigned and ensure the formation of the compounds 10a-j. The representative spectral data of 6-(4-amino-1-(2-(4-methylphenyl)-1-yl)benzofuran-3-yl)methyl-9-(1H-pyrazolo[3,4-d]pyrimidin-2-yl)benzo[d][1,2,3-b]oxazine (10j) is listed below.

¹H NMR (DMSO- d_6 , 400 MHz): δ ppm: 1.04 (m, 3H), 2.21 (s, 3H), 2.30-2.35 (m, 2H), 2.38-2.40 (m, 1H), 3.52 (m, 4H), 5.61 (s, 2H), 7.08-7.09 (m, 2H), 7.32-7.34 (d, J=8.7 Hz, 1H), 7.49-7.51 (d, J=8.7 Hz, 1H), 7.63-7.65 (m, 1H), 7.83-7.88 (d, J=7.2 Hz, 1H), 8.19 (s, 1H), 8.39 (s, 1H), 8.43 (s, 1H). **ESI-MS** (relative intensity): 508.1 (M+H)⁺ 100% (base peak).

¹³C NMR (DMSO- d_6 , 100 MHz): δ ppm: 1.04 (m, 3H), 2.21 (s, 3H), 2.30-2.35 (m, 2H), 2.38-2.40 (m, 1H), 3.52 (m, 4H), 5.61 (s, 2H), 7.08-7.09 (m, 2H), 7.32-7.34 (d, J=8.7 Hz, 1H), 7.49-7.51 (d, J=8.7 Hz, 1H), 7.63-7.65 (m, 1H), 7.83-7.88 (d, J=7.2 Hz, 1H), 8.19 (s, 1H), 8.39 (s, 1H), 8.43 (s, 1H).

***In vitro* Assay: PI3K ELISA Assay of p110 δ and p110 ϵ :**
Procedure: PI3K Kinase assay kit from Millipore was used to screen inhibitors of human PI3K-kinases p110 δ . The kinase reaction was setup with 25ng of the PI3K enzyme (3), 10nM PIP2 substrate & IX kinase reaction buffer in 25 μ l/well. Different conc. of the NCEs were added to the reaction mixture & incubated for 1h at 37°C. 25 μ l/well of Biotinylated-PIP3-EDTA working solution was added to the reaction well followed by 50 μ l of GRP1 working solution to all wells. The reaction was incubated for 1h at 20°C & washed 4 times with 200 μ l/well XTBST. SA-HRP working solution (50 μ l) is added & again incubated at 20°C for 1h. 100 μ l of the Substrate TMB is added to develop the color in the dark for 5-20 minutes and stopped by adding 100 μ l of the stop solution. The absorbance measured at 450nm & relative percentage of inhibition is calculated with respect to the positive control. *In vitro* PI3K δ (at 10, 100 & 1000 nM) and mTOR inhibitory activity (at 1000 nM) for test compounds (10a-j) are reported in Table 1.

Docking Study: As a representative compound from this series, molecular docking studies of 10j was carried out by using (Glide version 6.7) Schrödinger software to study an interaction of 10j with PI3K δ receptor. It was found that core benzofuran ring was oriented in the specificity pocket, N-methyl piperazine ring aligned towards hydrophobic region and substitution at R₁ position was found to be directed towards affinity pocket. Thus, docking results of 10j correlates with its potent *in vitro* PI3K δ activity.

Table 1: *In vitro* PI3K δ and mTOR inhibitory activity data of 10a-j

| S.No. | Structure | % inhibition PI3K δ at 3 conc. | | | % inhibition mTOR |
|---------|--------------|---------------------------------------|------|-------|-------------------|
| | | 1nM | 10nM | 100nM | |
| 10a* | | 68 | 90 | 102 | 15 |
| 10b* | | 70 | 88 | 100 | 12 |
| 10c* | | 55 | 78 | 98 | 16 |
| 10d | | 30 | 50 | 78 | 41 |
| 10e | | 10 | 45 | 53 | 18 |
| 10f | | 11 | 55 | 68 | 22 |
| 10g | | 40 | 58 | 78 | 38 |
| 10h* | | 69 | 90 | 99 | 13 |
| 10i* | | 70 | 95 | 108 | 11 |
| 10j* | | 90 | 98 | 112 | 10 |
| INK 654 | Std compound | 59 | 88 | 110 | 10 |

Summary and Conclusion:
Based on *in vitro* results, we can summarize that among ten novel compounds tested for PI3K δ and mTOR activities, compound 10a-c, 10h-j were found to be potent and selective PI3K δ inhibitors over mTOR.
In *in vitro* results validate our hypothesis of designing benzofuran based novel, potent and selective PI3K δ inhibitors as a bioisosteric replacement of quinoline ring of INK-654 with benzofuran ring system.
Further ADMET and *in vivo* profiling, followed by structure guided optimization of these lead molecules may likely to provide potent and PI3K δ selective inhibitor for the treatment of hematologic malignancies and inflammatory diseases.

Acknowledgement:
We are grateful to the management of Zydus Group for encouragement and the analytical department for providing analytical support.

References:
1. Nature chemical biology 6:2010(11):124.
2. J Biol Chem 280:10303-10310.
3. Bioorganic and Medicinal Chemistry Letter 12(2002):1687-1690.
4. Organic Synthetic. Col. Vol. 4(1963): p.590.
5. Bioorganic and Medicinal Chemistry Letter 18(2008): 5402-5405.

Chemistry: Overall, 10 compounds (10a-j) were prepared in good yield, under the mild reaction condition (Scheme 1) and were characterized with their physical, analytical and spectral data (1H NMR, ESI MS). The percentage yield in the final step was found to be in the range of 60-80%. The ESI MS and NMR spectral data of all the synthesized compounds were also found to be in conformity with the structures assigned and ensure the formation of the compounds 10a-j. The representative spectral data of 6-(4-amino-1-(2-(4-methylphenyl)-1-yl)benzofuran-3-yl)methyl-9-(1H-pyrazolo[3,4-d]pyrimidin-2-yl)benzo[d][1,2,3-b]oxazine (7a) is listed below.

¹H NMR (DMSO- d_6 , 400 MHz): δ ppm: 2.82 (s, 3H), 3.13-3.70 (m, 4H), 3.60-3.70 (m, 4H), 5.90 (s, 2H), 7.05 (s, 1H), 7.17-7.34 (m, 1H), 7.50-7.60 (m, 3H), 7.71-7.75 (m, 1H), 7.88-7.90 (m, 1H), 8.14-8.16 (m, 1H), 8.33 (s, 1H). **ESI-MS** (relative intensity): 522.2 (M+H)⁺ 100% (base peak).

¹³C NMR (DMSO- d_6 , 100 MHz): δ ppm: 2.82 (s, 3H), 3.13-3.70 (m, 4H), 3.60-3.70 (m, 4H), 5.90 (s, 2H), 7.05 (s, 1H), 7.17-7.34 (m, 1H), 7.50-7.60 (m, 3H), 7.71-7.75 (m, 1H), 7.88-7.90 (m, 1H), 8.14-8.16 (m, 1H), 8.33 (s, 1H).

***In vitro* Assay: PI3K ELISA Assay of p110 δ and p110 ϵ :**
Procedure: PI3K Kinase assay kit from Millipore was used to screen inhibitors of human PI3K-kinases p110 δ . The kinase reaction was setup with 25ng of the PI3K enzyme (3), 10nM PIP2 substrate & IX kinase reaction buffer in 25 μ l/well. Different conc. of the NCEs were added to the reaction mixture & incubated for 1h at 37°C. 25 μ l/well of Biotinylated-PIP3-EDTA working solution was added to the reaction well followed by 50 μ l of GRP1 working solution to all wells. The reaction was incubated for 1h at 20°C & washed 4 times with 200 μ l/well XTBST. SA-HRP working solution (50 μ l) is added & again incubated at 20°C for 1h. 100 μ l of the Substrate TMB is added to develop the color in the dark for 5-20 minutes and stopped by adding 100 μ l of the stop solution. The absorbance measured at 450nm & relative percentage of inhibition is calculated with respect to the positive control. *In vitro* PI3K δ (at 10, 100 & 1000 nM) and mTOR inhibitory activity (at 1000 nM) for test compounds (7a-j) are reported in Table 1.

Docking Study: As a representative compound from this series, molecular docking studies of 7a was carried out by using (Glide version 6.7) Schrödinger software to study an interaction of 7a with PI3K δ receptor (PDB ID: 2W4K). It was found that core Quinoline ring was oriented in the specificity pocket, N-methyl piperazine ring aligned towards hydrophobic region and substitution at R₁ position was found to be directed towards affinity pocket. Thus, docking results of 7a correlates with its potent *in vitro* PI3K δ activity.

Table 1: *In vitro* PI3K δ and mTOR inhibitory activity data of 7a-j

| S.No. | Structure | % inhibition PI3K δ at 3 conc. | | | % inhibition mTOR |
|---------|--------------------|---------------------------------------|------|-------|-------------------|
| | | 1nM | 10nM | 100nM | |
| 7a* | | 66 | 82 | 110 | 10 |
| 7b* | | 62 | 84 | 105 | 12 |
| 7c* | | 50 | 75 | 94 | 14 |
| 7d | | 45 | 52 | 60 | 30 |
| 7e | | 35 | 44 | 52 | 35 |
| 7f | | 42 | 58 | 70 | 28 |
| 7g | | 32 | 54 | 70 | 35 |
| 7h* | | 70 | 92 | 108 | 10 |
| 7i* | | 72 | 95 | 105 | 12 |
| 7j* | | 68 | 94 | 98 | 15 |
| INK 654 | Reference Compound | 59 | 88 | 110 | 10 |

Chemistry: Overall, 10 compounds (7a-j) were prepared in good yield, under the mild reaction condition (Scheme 1) and were characterized with their physical, analytical and spectral data (1H NMR, ESI MS). The percentage yield in the final step was found to be in the range of 60-80%. The ESI MS and NMR spectral data of all the synthesized compounds were also found to be in conformity with the structures assigned and ensure the formation of the compounds 7a-j. The representative spectral data of 6-(4-amino-1-(2-(4-methylphenyl)-1-yl)benzofuran-3-yl)methyl-9-(1H-pyrazolo[3,4-d]pyrimidin-2-yl)benzo[d][1,2,3-b]oxazine (7a) is listed below.

¹H NMR (DMSO- d_6 , 400 MHz): δ ppm: 2.82 (s, 3H), 3.13-3.70 (m, 4H), 3.60-3.70 (m, 4H), 5.90 (s, 2H), 7.05 (s, 1H), 7.17-7.34 (m, 1H), 7.50-7.60 (m, 3H), 7.71-7.75 (m, 1H), 7.88-7.90 (m, 1H), 8.14-8.16 (m, 1H), 8.33 (s, 1H). **ESI-MS** (relative intensity): 522.2 (M+H)⁺ 100% (base peak).

¹³C NMR (DMSO- d_6 , 100 MHz): δ ppm: 2.82 (s, 3H), 3.13-3.70 (m, 4H), 3.60-3.70 (m, 4H), 5.90 (s, 2H), 7.05 (s, 1H), 7.17-7.34 (m, 1H), 7.50-7.60 (m, 3H), 7.71-7.75 (m, 1H), 7.88-7.90 (m, 1H), 8.14-8.16 (m, 1H), 8.33 (s, 1H).

***In vitro* Assay: PI3K ELISA Assay of p110 δ and p110 ϵ :**
Procedure: PI3K Kinase assay kit from Millipore was used to screen inhibitors of human PI3K-kinases p110 δ . The kinase reaction was setup with 25ng of the PI3K enzyme (3), 10nM PIP2 substrate & IX kinase reaction buffer in 25 μ l/well. Different conc. of the NCEs were added to the reaction mixture & incubated for 1h at 37°C. 25 μ l/well of Biotinylated-PIP3-EDTA working solution was added to the reaction well followed by 50 μ l of GRP1 working solution to all wells. The reaction was incubated for 1h at 20°C & washed 4 times with 200 μ l/well XTBST. SA-HRP working solution (50 μ l) is added & again incubated at 20°C for 1h. 100 μ l of the Substrate TMB is added to develop the color in the dark for 5-20 minutes and stopped by adding 100 μ l of the stop solution. The absorbance measured at 450nm & relative percentage of inhibition is calculated with respect to the positive control. *In vitro* PI3K δ (at 10, 100 & 1000 nM) and mTOR inhibitory activity (at 1000 nM) for test compounds (7a-j) are reported in Table 1.

Docking Study: As a representative compound from this series, molecular docking studies of 7a was carried out by using (Glide version 6.7) Schrödinger software to study an interaction of 7a with PI3K δ receptor (PDB ID: 2W4K). It was found that core Quinoline ring was oriented in the specificity pocket, N-methyl piperazine ring aligned towards hydrophobic region and substitution at R₁ position was found to be directed towards affinity pocket. Thus, docking results of 7a correlates with its potent *in vitro* PI3K δ activity.

Table 1: *In vitro* PI3K δ and mTOR inhibitory activity data of 7a-j

| S.No. | Structure | % inhibition PI3K δ at 3 conc. | | | % inhibition mTOR |
|---------|--------------------|---------------------------------------|------|-------|-------------------|
| | | 1nM | 10nM | 100nM | |
| 7a* | | 66 | 82 | 110 | 10 |
| 7b* | | 62 | 84 | 105 | 12 |
| 7c* | | 50 | 75 | 94 | 14 |
| 7d | | 45 | 52 | 60 | 30 |
| 7e | | 35 | 44 | 52 | 35 |
| 7f | | 42 | 58 | 70 | 28 |
| 7g | | 32 | 54 | 70 | 35 |
| 7h* | | 70 | 92 | 108 | 10 |
| 7i* | | 72 | 95 | 105 | 12 |
| 7j* | | 68 | 94 | 98 | 15 |
| INK 654 | Reference Compound | 59 | 88 | 110 | 10 |

Chemistry: Overall, 10 compounds (7a-j) were prepared in good yield, under the mild reaction condition (Scheme 1) and were characterized with their physical, analytical and spectral data (1H NMR, ESI MS). The percentage yield in the final step was found to be in the range of 60-80%. The ESI MS and NMR spectral data of all the synthesized compounds were also found to be in conformity with the structures assigned and ensure the formation of the compounds 7a-j. The representative spectral data of 6-(4-amino-1-(2-(4-methylphenyl)-1-yl)benzofuran-3-yl)methyl-9-(1H-pyrazolo[3,4-d]pyrimidin-2-yl)benzo[d][1,2,3-b]oxazine (7a) is listed below.

¹H NMR (DMSO- d_6 , 400 MHz): δ ppm: 2.82 (s, 3H), 3.13-3.70 (m, 4H), 3.60-3.70 (m, 4H), 5.90 (s, 2H), 7.05 (s, 1H), 7.17-7.34 (m, 1H), 7.50-7.60 (m, 3H), 7.71-7.75 (m, 1H), 7.88-7.90 (m, 1H), 8.14-8.16 (m, 1H), 8.33 (s, 1H). **ESI-MS** (relative intensity): 522.2 (M+H)⁺ 100% (base peak).

¹³C NMR (DMSO- d_6 , 100 MHz): δ ppm: 2.82 (s, 3H), 3.13-3.70 (m, 4H), 3.60-3.70 (m, 4H), 5.90 (s, 2H), 7.05 (s, 1H), 7.17-7.34 (m, 1H), 7.50-7.60 (m, 3H), 7.71-7.75 (m, 1H), 7.88-7.90 (m, 1H), 8.14-8.16 (m, 1H), 8.33 (s, 1H).

***In vitro* Assay: PI3K ELISA Assay of p110 δ and p110 ϵ :**
Procedure: PI3K Kinase assay kit from Millipore was used to screen inhibitors of human PI3K-kinases p110 δ . The kinase reaction was setup with 25ng of the PI3K enzyme (3), 10nM PIP2 substrate & IX kinase reaction buffer in 25 μ l/well. Different conc. of the NCEs were added to the reaction mixture & incubated for 1h at 37°C. 25 μ l/well of Biotinylated-PIP3-EDTA working solution was added to the reaction well followed by 50 μ l of GRP1 working solution to all wells. The reaction was incubated for 1h at 20°C & washed 4 times with 200 μ l/well XTBST. SA-HRP working solution (50 μ l) is added & again incubated at 20°C for 1h. 100 μ l of the Substrate TMB is added to develop the color in the dark for 5-20 minutes and stopped by adding 100 μ l of the stop solution. The absorbance measured at 450nm & relative percentage of inhibition is calculated with respect to the positive control. *In vitro* PI3K δ (at 10, 100 & 1000 nM) and mTOR inhibitory activity (at 1000 nM) for test compounds (7a-j) are reported in Table 1.

Docking Study: As a representative compound from this series, molecular docking studies of 7a was carried out by using (Glide version 6.7) Schrödinger software to study an interaction of 7a with PI3K δ receptor (PDB ID: 2W4K). It was found that core Quinoline ring was oriented in the specificity pocket, N-methyl piperazine ring aligned towards hydrophobic region and substitution at R₁ position was found to be directed towards affinity pocket. Thus, docking results of 7a correlates with its potent *in vitro* PI3K δ activity.

Table 1: *In vitro* PI3K δ and mTOR inhibitory activity data of 7a-j

| S.No. | Structure | % inhibition PI3K δ at 3 conc. | | | % inhibition mTOR |
|---------|--------------------|---------------------------------------|------|-------|-------------------|
| | | 1nM | 10nM | 100nM | |
| 7a* | | 66 | 82 | 110 | 10 |
| 7b* | | 62 | 84 | 105 | 12 |
| 7c* | | 50 | 75 | 94 | 14 |
| 7d | | 45 | 52 | 60 | 30 |
| 7e | | 35 | 44 | 52 | 35 |
| 7f | | 42 | 58 | 70 | 28 |
| 7g | | 32 | 54 | 70 | 35 |
| 7h* | | 70 | 92 | 108 | 10 |
| 7i* | | 72 | 95 | 105 | 12 |
| 7j* | | 68 | 94 | 98 | 15 |
| INK 654 | Reference Compound | 59 | 88 | 110 | 10 |

Chemistry: Overall, 10 compounds (7a-j) were prepared in good yield, under the mild reaction condition (Scheme 1) and were characterized with their physical, analytical and spectral data (1H NMR, ESI MS). The percentage yield in the final step was found to be in the range of 60-80%. The ESI MS and NMR spectral data of all the synthesized compounds were also found to be in conformity with the structures assigned and ensure the formation of the compounds 7a-j. The representative spectral data of 6-(4-amino-1-(2-(4-methylphenyl)-1-yl)benzofuran-3-yl)methyl-9-(1H-pyrazolo[3,4-d]pyrimidin-2-yl)benzo[d][1,2,3-b]oxazine (7a) is listed below.

¹H NMR (DMSO- d_6 , 400 MHz): δ ppm: 2.82 (s, 3H), 3.13-3.70 (m, 4H), 3.60-3.70 (m, 4H), 5.90 (s, 2H), 7.05 (s, 1H), 7.17-7.34 (m, 1H), 7.50-7.60 (m, 3H), 7.71-7.75 (m, 1H), 7.88-7.90 (m, 1H), 8.14-8.16 (m, 1H), 8.33 (s, 1H). **ESI-MS** (relative intensity): 522.2 (M+H)⁺ 100% (base peak).

¹³C NMR (DMSO- d_6 , 100 MHz): δ ppm: 2.82 (s, 3H), 3.13-3.70 (m, 4H), 3.60-3.70 (m, 4H), 5.90 (s, 2H), 7.05 (s, 1H), 7.17-7.34 (m, 1H), 7.50-7.60 (m, 3H), 7.71-7.75 (m, 1H), 7.88-7.90 (m, 1H), 8.14-8.16 (m, 1H), 8.33 (s, 1H).

***In vitro* Assay: PI3K ELISA Assay of p110 δ and p110 ϵ :**
Procedure: PI3K Kinase assay kit from Millipore was used to screen inhibitors of human PI3K-kinases p110 δ . The kinase reaction was setup with 25ng of the PI3K enzyme (3), 10nM PIP2 substrate & IX kinase reaction buffer in 25 μ l/well. Different conc. of the NCEs were added to the reaction mixture & incubated for 1h at 37°C. 25 μ l/well of Biotinylated-PIP3-EDTA working solution was added to the reaction well followed by 50 μ l of GRP1 working solution to all wells. The reaction was incubated for 1h at 20°C & washed 4 times with 200 μ l/well XTBST. SA-HRP working solution (50 μ l) is added & again incubated at 20°C for 1h. 100 μ l of the Substrate TMB is added to develop the color in the dark for 5-20 minutes and stopped by adding 100 μ l of the stop solution. The absorbance measured at 450nm & relative percentage of inhibition is calculated with respect to the positive control. *In vitro* PI3K δ (at 10, 100 & 1000 nM) and mTOR inhibitory activity (at 1000 nM) for test compounds (7a-j) are reported in Table 1.

Docking Study: As a representative compound from this series, molecular docking studies of 7a was carried out by using (Glide version 6.7) Schrödinger software to study

