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Synthesis and Biological Evaluation of 4-(Piperid-3-yl)amino Substituted 6-Pyridylquinazolines as Potent PI3Kδ Inhibitors

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Abstract: PI3Kδ is an intriguing target for developing anti-cancer agent. In this study, a new series of 4-(piperid-3-yl)amino substituted 6-pyridylquinazoline derivatives were synthesized. After biological evaluation, compounds **A5** and **A8** were identified as potent PI3Kδ inhibitors, with IC₅₀ values of 1.3 and 0.7 nM, respectively, which are equivalent to or better than idelalisib (IC₅₀ = 1.2 nM). Further PI3K isoforms selectivity evaluation showed that compound **A5** afforded excellent PI3Kδ selectivity over PI3Kα, PI3Kβ and PI3Kγ. **A8** exhibited superior PI3Kδ/γ selectivity over PI3Kα and PI3Kβ. Moreover, compounds **A5** and **A8** selectively exhibited anti-proliferation against SU-DHL-6 *in vitro* with IC₅₀ values of 0.16 and 0.12 μM. Western blot analysis indicated that **A8** formed three key H-bonds action with PI3Kδ, which may account for its potent inhibition of PI3Kδ. These findings indicate that 4-(piperid-3-yl)amino substituted 6-pyridylquinazoline derivatives were potent PI3Kδ inhibitors with distinctive PI3K-isoforms and anti-proliferation profiles.

Keywords: PI3Kδ inhibitors; PI3K inhibitors; 6-pyridylquinazolines; anti-proliferation.

1. Introduction

Phosphoinositide 3-kinases (PI3Ks) are members of a family of key intracellular enzymes. PI3Ks mediate the phosporylation procedure, and trigger the downstream signal transducer AKT (protein kinase B) and mTOR (Mannalian target of rapamycin) to regulate cellular activities such as cell proliferation, differentiation, migration and survival. PI3K family consists of three classes. The class I PI3Ks are well characterized, which are further individed into PI3K α , β , δ and γ .¹ PI3K α and β are widely expressed, whereas PI3K δ and γ are predominantly restricted in hematopoietic cells.² Consequently, inhibition of PI3K δ (or mixed PI3K δ/γ) was thought to enable to combat cancers, immunological and inflammatory conditions and development of small molecule PI3K δ inhibitors receives significant interest from industrial and academic community.³

Many classes of PI3K\delta inhibitors have been investigated, and some have been approved or entered into clinical trials for treatment of leukemia, lymphoma, asthma, chronic obstructive pulmonary disease (COPD), sjogren's syndrome, activated PI3Kδ syndrome (APDS), and rheumatoid arthritis.⁴ The first-in-class orally potent selective PI3K δ inhibitor idelalisib (1) was approved by FDA (US Food and Drug Administration) in 2014 for treating chronic lymphocytic leukemia (CLL), follicular lymphoma (FL) and small lymphocytic lymphoma (SLL).⁵ Although idelalisib gave significant efficacy in clinic, it showed severe toxicity leading to its approval with black-box warning. ⁶ Later in 2018, the selective PI3K δ/γ inhibitors duvelisib (2) was approved by FDA also for treating CLL, FL and SLL, which was structurally similar to idelalisib.⁷ Duvelisib was reported to show lower side effects than idelalisib. Umbralisib (3) was another potent selective PI3K\delta inhibitor, studied in phase III clinical trials for its efficacy against B-cell lymphoma.⁸ Leniolisib (4), developed by Novartis, was an orally potent selective PI3Kδ inhibitor which recently entered phase III clinical evaluation for rare autoimmune diseases such as APDS and Sjogren's syndrome.^{9, 10} Nemiralisib (5), developed by GSK (GlaxoSmithKline), was in phase II trials for treatment of asthma and COPD ¹¹ (Figure 1). Overall, these approved and late-stage clinical selective PI3K\delta inhibitors are categorized into two classes and exemplified as follows: (1) propeller-shaped inhibitors such as idelalisib, duvelisib, and umbralisib, (2) flat-shaped inhibitors such as leniolisib and nemiralisib.¹² Considering the reported propeller-shaped PI3Kδ inhibitors

demonstrating severe side effects, there is greater need to develop new classes of PI3K δ inhibitors differentiated from propeller-shaped ones, so as to investigate the medication regimen and improve the safety.¹³ Herein, we reported our drug effort on development of PI3K δ inhibitors which differed from propeller-shaped ones.



Figure 1. Representative structures of approved and late-stage clinical PI3Kδ inhibitors.

Recently, we disclosed a series of potential PI3K\delta inhibitors containing a quinazoline scaffold which were significantly different from the propeller-shaped inhibitors.¹⁴ Exploring the 4-substitutents of 6-pyridylquinazoline led to the 4-aniline-6-pyridylquinazolines exemplified by 6 (PI3K δ : IC₅₀ = 9.3 nM) and the 4-pyrrolidineamino-6-pyridylquinazolines exemplified by 7 (PI3K δ : IC₅₀ = 2.7 nM).^{15, 16} Simply switching the 4-pyrrolidineamino to 4-pyrrolidineoxy (or 4-(piperid-3-yl)amino) group made the PI3K δ inhibitory activity retained or enhanced, such as 8 (PI3K δ : IC₅₀ = 4.9 nM) and 9 (PI3K δ : IC₅₀ = 3.0 nM).¹⁷ In particular, we speculated that larger 4-(piperid-3-yl)amino moiety had a bigger twist angle compared to the 4-pyrrolidineamino motif linked to 6-pyridylquinazoline flat, which may contribute to

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balance the PI3K δ -selectivity (**Figure 2**). Therefore, we carry out a further structural investigation by introducing H-bond acceptor groups such as cyano, nitro and trifluoromethyl at 5-position of 6-pyridyl moiety and several side chains such as hydrophilic and hydrophobic groups as the tails linked to the 4-(piperid-3-yl)amino moiety, so as to establish a comprehensive structure-activity relationship (SAR) and the bioactive profiles. We now report the synthesis and biological evaluation of this series of new 4-(piperid-3-yl)amino substituted 6-pyridylquinazoline analogues as potent PI3K δ inhibitors, notably, of which several compounds showed significant selectivity in both PI3K δ inhibition and anti-proliferative activity of cells lines (**Figure 3**).

(a)



compound7compound 9Figure 2. (a) Chemical structures of 6, 7, 8 and 9; (b) 3D structures of 7 and 9.



R¹ = H-bond acceptor groups (CN, NO₂, CF₃) R = hydrophilic and hydrophobic groups

Figure 3. Current design of 4-(piperid-3-yl)amino substituted 6-pyridylquinazolines as PI3Kδ inhibitors.

2. Results and discussion

2.1 Chemistry

The synthetic route of theses 4-(piperid-3-yl)amino substituted quinazoline derivatives A1-14 was outlined in Scheme 1. 6-Bromo-4-chloroquinazoline (10) reacted with (S)-1-Boc-3-aminopiperidine in the presence of N, N-diisopropylethylamine (DIPEA) in refluxing i-PrOH gave intermediate 11. Subsequently, R¹-substituted 2-methoxy-3-bromopyridine reacted with bis(pinacolato)diboron under the Miyaura borvlation condition without further separation to produce R¹-substituted which were immediately treated with **11** using 6-methoxypyridinylboronate, Pd-catalysized Suzuki-coupling reaction to provide 12a-c in one pot. Removal of Boc-group of **12a-c** by TFA at room temperature afforded **13a-c**. **13a** was acylated with varied acids to generate compounds A1-A9. Compounds A10 and A12 were prepared following acylation procedure by TFAA, and compounds A11, A13 and A14 were prepared following nucleophilic substitution procedure by ethyl bromoacetate and bromoethanol, respectively (Scheme 1).



Scheme 1. Reagents and conditions: (a) (S)-1-Boc-3-aminopiperidine, DIPEA, i-PrOH, reflux, 85%; (b) (i) R¹-substituted 3-bromo-6-methoxypyridine KOAc, PdCl₂(dppf), dioxane, reflux, 2.5 h; (ii) R¹-substituted 6-methoxypyridinylboronate, Na₂CO₃, PdCl₂(dppf), dioxane/H₂O, reflux, 2.5 h; 57-79 % for two steps; (c) CH₂Cl₂:TFA = 5:1, rt, 2 h, 23-91 %; (d) diverse acids, DMF, HATU, DIPEA, rt, 12 h, 34-73 %; (e) TFAA, DMF, DIPEA, rt, 55 % for A10 and 57% for A12; (f) BrCH₂CO₂Et, CH₂Cl₂, DIPEA, reflux, 43 % for A11 and 57 % for A13; (g) BrCH₂CH₂OH, CH₃CN, DIPEA, reflux, 23 %.

2.2 PI3Kô inhibitory activities and structure-activity relationship (SAR) study

All the newly synthesized 4-(piperid-3-yl)amino substituted 6-pyridylquinazoline analogues were firstly assessed for their inhibitory activities against PI3K δ . Idelalisib was employed as the positive control. The IC₅₀ values were illustrated in Table 1. As expected, all the 4-(piperid-3-yl)amino substituted 6-pyridylquinazolines showed significantly potent PI3K δ inhibitory activities with IC₅₀ values < 100 nM, suggesting the 4-(piperid-3-yl)amino-6-(6-methoxylpyridin-3-yl)quinazoline moiety is critical for maintaining the PI3K δ inhibition. Based on the previously observed SAR from 4-pyrrolidineamino-6-6-pyridylquinazoline series, the cyano group was firstly fixed at the 5-position of 6-methoxylpyridin-3-yl moiety, several R side chains including hydrophilic and hydrophobic group were investigated. Compound **A1** bearing a hydrophilic tetrahydro-2*H*-pyran-4-carbonyl tail showed potent PI3K δ inhibitory activity with an IC₅₀ value of 7.7 nM, while replacement of the tetrahydro-2*H*-pyranyl group with 4-methylpiperidyl group gave equivalent potency (A2, $IC_{50} = 9.4$ nM). Removal of the methyl group led to compound **12c** showing 4-fold improved potency with IC_{50} value of 2.5 nM, whereas replacement with the 2-aminopropanoyl group retained approximately potent PI3K δ activity (A4, IC₅₀ = 3.7 nM). Tuning R to *N*,*N*-dimethylaminoacetyl group resulted in improving PI3K δ inhibitory activity (A5: IC₅₀ = 1.3 nM) which was equal to idelalisib (IC₅₀ = 1.2 nM). Subsequently, compound A6 bearing a hydrophobic chloroacetyl side chain afforded an IC₅₀ value of 8.9 nM, while compound A7 bearing bromopropional tail gave lower PI3K δ inhibition with an IC₅₀ value of 14.3 nM. Moreover, the heteroaromatic tails were attempted. Compound A8 with a nicotinic terminal showed the most potent PI3Kδ inhibitory activity with an IC₅₀ value of 0.7 nM, two-fold higher than idelalisib. Introduction of the 5-nitro-2-furoyl group gave compound A9 showing PI3K δ IC₅₀ value of 3.9 nM. Furthermore, two hydrophobic terminal group were synthesized leading to compounds A10 and A11 tailed trifluoroacetyl and ethoxylcarbonylmethyl group, respectively, which displayed comparable potency, with IC_{50} value of both 3.2 nM, whereas further changing the cyano group into nitro (A12, $IC_{50} = 27 \text{ nM}$) and trifluoromethyl group (A13, $IC_{50} = 6.8 \text{ nM}$) on 6-pyridinyl moiety led to the potency 9- and 2-fold reduced, respectively. An attempt to introduce the hydroxyethyl tail led to compound A14 showed PI3K δ inhibition retainable, with an IC₅₀ value of 4.7 nM. Overall, in this first PI3K δ inhibition evaluation, two compounds A5 and A8 showed IC₅₀ values of < 2 nM, which was equivalent to or better than idelalisib, and six compounds including A3, A4, A9, A10, A11 and A14 demonstrated IC_{50} values between 2 and 5 nM, which was comparable to idelalisib.

All the target compounds were subsequently examined for their cellular anti-proliferative activity against PI3K δ -susceptible human malignant B-cell line SU-DHL-6. The results were summarized in Table 1. In general, except for compound A12 with weak anti-proliferative efficacy against SU-DHL-6, the other compounds showed favorable anti-proliferation with micromolar IC₅₀ values or below. Both of compounds A5 and A8, the most potent PI3K δ inhibitors, exhibited significantly anti-proliferative activity against SU-DHL-6 cell line with IC₅₀ values of 0.16 and 0.12 μ M, respectively.

Table 1. PI3K δ inhibitory activity and SAR study of 4-(piperid-3-yl)amino substituted quinazolines ^a



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Compds	\mathbf{R}^1	R	ΡΙ3Κδ	SU-DHL
compus	i i i i i i i i i i i i i i i i i i i	i c	IC ₅₀ (nM)	IC_{50} (μM)
A1	CN		7.7 ± 2.1	0.15 ± 0.01
A2	CN	- z ^z O	9.4 ± 0.1	0.42 ± 0.06
A3	CN	r de la constantina de la cons	2.5 ± 0.3	2.44 ± 0.43
A4	CN	NH2 ONH2	3.7 ± 0.2	0.44 ± 0.02
A5	CN	^{2²} − N −	1.3 ± 0.4	0.16 ± 0.05
A6	CN	CI O	8.9 ± 1.0	0.38 ± 0.14
A7	CN	ζζ⁵→ Br O	14.3 ± 1.0	0.015 ± 0.00
A8	CN	P C O	0.7 ± 0.1	0.12 ± 0.05
A9	CN		3.9 ± 0.7	0.14 ± 0.02
A10	CN	v ^{z^s} ↓CF ₃ O	3.2 ± 0.2	0.21 ± 0.02
A11	CN	×35 0	3.2 ± 0.9	0.50 ± 0.12
A12	NO ₂	Č ⁵ OCF₃ O	27.0 ± 3.1	127 ± 32

A13	CF ₃	, est of the second sec	6.8 ± 1.8	0.23 ± 0.05
A14	CN). Contraction of the second s	4.7 ± 2.1	2.05 ± 0.32
Idelalisib	-	-	1.2 ± 0.2	0.033 ± 0.10

^a The IC₅₀ values are shown as the mean \pm SD (nM) from two separate experiments.

2.3 Isoform selectivity of the new PI3Kô inhibitors

To further evaluated the selectivity of the title compounds, four compounds A4, A5, **A8**, and **A9** were evaluated for their inhibitory activities against a panel of PI3K kinases. The data were listed in Table 2. Obviously, compounds A4 and A5 were selective PI3K δ inhibitors which showed weak inhibition against PI3K β and PI3K γ , and moderate inhibition against PI3Ka. In particular, A5 displayed 17-fold, 115-fold and 192-fold PI3K δ selectivity over PI3K α , PI3K β and PI3K γ , respectively. A8 was a PI3K γ/δ inhibitor with IC_{50} values of 2.0 and 0.7 nM against PI3K γ and PI3K δ , respectively. Notably, A8 exhibited 170-fold and 13-fold selectivity of PI3K δ over PI3K α and PI3K β , respectively. A9 was a PI3K α/δ inhibitor with IC₅₀ values of 1.8 and 3.9 nM against PI3K α and PI3K δ , respectively, and showed moderate inhibition against PI3K β and PI3Ky. Compounds selectively targeting PI3K δ were thought to be capable of attenuating side effects. However, the orally potent PI3K δ inhibitor idelalisib afforded severe toxicity in the clinical validation but the approved pan-PI3K inhibitor copanlisib demonstrated lower toxicity and fewer side effects compared to idelalisib. Therefore, the PI3K isoforms selectivity is still an open question. In this series, particularly, compound A5 was a selective PI3K δ inhibitor, and A8 was a PI3K γ/δ inhibitor (Table 2).

compds	IC ₅₀ (nM)				
	ΡΙ3Κα	ΡΙ3Κβ	ΡΙ3Κγ	ΡΙ3Κδ	
A4	18.2 ± 1.4	147.4 ± 9.5	210.7 ± 7.4	3.7 ± 0.2	
A5	21.8 ± 2.1	150.1 ± 5.4	250.4 ± 8.7	1.3 ± 0.4	
A8	119.5 ± 6.2	9.1 ± 1.2	2.0 ± 0.2	0.7 ± 0.1	
A9	1.8 ± 0.3	31.4 ± 2.1	19.3 ± 2.1	3.9 ± 0.7	
Idelalisib	177.2 ± 4.4	139.1 ± 5.2	145.3 ± 7.9	1.2 ± 0.2	

Table 2. PI3K isoform selectivity of compounds against Class I PI3Ks ^a

^a The IC₅₀ values are shown as the mean \pm SD (nM) from two separate experiments.

2.4 Cellular anti-proliferation evaluation

Either the pan-PI3K or selective PI3K δ (or PI3K δ/γ) inhibitor mainly aimed to treat cancers. Consequently, furthermore, four compounds A4, A5, A8, and A9 were tested for their anti-proliferative activities against five cancer lines including B cell lines SU-DHL-6, Ramos, Raji, and solid cancer lines HCT-116 and MCF-7 *in vitro*. Idelalisib and BEZ235 (a clinical PI3K/mTOR dual inhibitor) were used as positive control drugs. The data were shown in Table 3. It was found four compounds as well as idelalisib showed significantly selective anti-proliferative activity. They were effective only against SU-DHL-6 but no efficacy against Ramos, Raji, HCT-116 and MCF-7. Contrarily, BEZ235 exhibited potent anti-proliferation against all five cell lines. In a word, four compounds A4, A5, A8, and A9 showed similar anti-proliferative profiles to idelalisib, which remarkably differed from PI3K/mTOR dual inhibitor BEZ235.

aamnaunda	IC ₅₀ (μM)				
compounds	SU-DHL-6	Ramos	Raji	HCT-116	MCF-7
A4	0.44 ± 0.02	>10	>10	>10	>10
A5	0.16 ± 0.05	>10	>10	>10	>10
A8	0.12 ± 0.05	>10	>10	>10	>10
A9	0.14 ± 0.02	>10	>10	2.12 ± 0.97	>10
Idelalisib	0.033 ± 0.10	>10	>10	>10	>10
BEZ235	-	0.40 ± 0.00	0.52 ± 0.20	0.30 ± 0.12	0.2 ± 0.02

Table 3. Anti-proliferative activities of selected compounds in vitro^a

^a The IC₅₀ values are average \pm SD (nM) of at least three independent experiments in triplicate.

2.5 PI3K_δ pathway cascade evaluation by AKT phosphorylation

Compound **A8** was a selective PI3K δ inhibitor, which was further investigated for its effects on PI3K-mediated signaling in SU-DHL-6 cell line by western blot analysis. As shown in Figure 4, compound **A8** displayed good reduction of phosphorylation of AKT (Ser473) with a concentration-dependent manner. This result indicated that compound **A8** enable to inhibit the PI3K/AKT/mTOR signaling pathway (Figure 4).



Figure 4. Effect of compound **A8** on AKT and p-AKT⁴⁷³. SU-DHL-6 cells were treated with various concentration of **A8** for 8h and western blot for indicated proteins was performed. β -actin was used as a loading control.

2.6 Molecular modeling study

The molecular docking analysis of compound **A8** within human PI3K δ enzyme were carried out to understand the potent PI3K δ inhibition. As shown in Figure 5, the docked pose of **A8** showed the favorable interactions within the PI3K δ binding pocket (PDB code: 6G6W). The quinazoline scaffold was engaged in H-bond contacts with Val828 in the hinge region, while the cyano group linked to 6-pyridyl moiety formed another H-bond with Lys779 in the affinity pocket. Moreover, the 4-(piperid-3-yl)amino side chain extended to solvent region and the carbonyl group generated an H-bond with Asn836. This region was close to the PI3K δ isoform specificity pocket and had a hydrogen bond with Asn836 which were key factors to generate specific PI3K δ selectivity ¹⁵. After all, these results may account for the potent PI3K δ inhibition for **A8** (Figure 5).



Figure 5. Molecular docking studies of compound A8 into the site of PI3K δ (PDB code: 6G6W). Compound is shown as sticks. H-bonds within 2.5 Å are shown as yellow dashed lines.

3. Conclusion

In summary, we described the design, synthesis and biological evaluation of a novel series of 4-(piperid-3-yl)amino substituted 6-pyridylquinazoline derivatives as PI3Kδ inhibitors. The structure-activity relationship was discussed and many derivatives showed nanomolar PI3K\delta inhibitory activities. Compounds A5 and A8 displayed significantly potent PI3K δ inhibitory activities with IC₅₀ values of 1.3 and 0.7 nM, respectively, which was equivalent to or better than idelalisib ($IC_{50} = 1.2$ nM). Compounds A5 and A8 showed highly efficacious anti-proliferation against PI3Kδ-sensitive human malignant B-cell line SU-DHL-6 in vitro with IC₅₀ values of 0.16 and 0.12 µM. Compounds A5 showed superior isoform selectivity over PI3K α , PI3K β and PI3K γ , and A8 showed favorable PI3K γ/δ selectivity over PI3K α and PI3K β . Interestingly, both A5 and A8 exhibited similarly potent anti-proliferation profiles to idelalisib. Western blot analysis of **A8**-treated SU-DHL-6 cell line indicated that **A8** reduced the AKT^{S473} phosphorylation. Molecular docking simulation suggested that the high potency of A8 was attributed to the formation of three key H-bonds. These findings indicate the 4-(piperid-3-yl)amino substituted 6-pyridylquinazoline derivatives were potent PI3K δ inhibitors with distinctive PI3K-isoforms and anti-proliferation profiles.

4. Experimental section

4.1 General methods

¹H NMR (400 MHz) and ¹³C NMR (100 MHz) spectra were recorded on a 400 Bruker NMR spectrometer with TMS as an internal reference. Mass spectra (MS) were routinely recorded on an API 3200 LC/MS spectrometer. High-resolution MS (HRMS) were measured by electrospray ionization (positive mode) on a quadrupole time-of-flight (QTOF) mass spectrometer (Maxis Q-TOF, Bruker Inc.). The reagents and solvents were purchased commercially and used without further purification unless specially described otherwise. The starting material 6-bromo-4-chloroquinazoline (**10**) was prepared according to our previously reported methods ¹⁵.

4.1.1 (S)-tert-Butyl 3-((6-bromoquinazolin-4-yl) amino) piperidine-1-carboxylate (11)

To a solution of (S)-1-Boc-3-aminopiperidine (6.20 g, 30.99 mmol) in iso-propanol (20 mL), was slowly added 6-bromo-4-chloroquinazoline (5.00 g, 20.66 mmol) and DIPEA (5.33 g, 41.32mmol). The mixture was stirred at 85°C for 2.5 h, and cooled to room temperature. After the saturated NaCl solution was added, the reaction mixture was extracted by EtOAc. Dried over anhydrous Na₂SO₄, the liquid was concentrated and purified by silica gel chromatography (CH₂Cl₂/CH₃OH, 100:1) to produce compound **11** (7.15 g, 85 %) as a yellow solid. ¹H-NMR (400 MHz, CDCl₃) δ 8.66 (s, 1H, ArH), 7.98 (s, 1H, ArH), 7.89-7.79 (m, 2H, ArH), 6.98 (s, 1H, NH), 4.41-4.39 (m, 1H, CH), 3.82-3.32 (m, 4H, 2×CH₂), 2.45-2.28 (m, 2H, CH₂), 1.98-1.75 (m, 2H, CH₂), 1.48 (s, 9H, 3×CH₃) ppm. MS (ESI) m/z: [M+H]⁺ =406.9, 408.9.

4.1.2 (S)-tert-Butyl 3-((6-(6-methoxypyridin-3-yl)quinazolin-4-yl) amino) piperidine-1-carboxylate (12a)

Compound **12a** was prepared according our reported method. To a solution mixture of **11** (1.20 g, 2.95mmol) in DME/H₂O (8 mL/2 mL), were added the 2-methoxy-5-(4,4,5,5-tetramethyl-1,3,2-dioxaborolan-2-yl)nicotinonitrile (1.01 g, 3.9 mmol), PdCl₂(dppf) (270 mg, 0.12 mmol) and Na₂CO₃ (636 mg, 6.0 mmol). The mixture was refluxed under argon atmosphere for 4 h, cooled to room temperature and concentrated. The residue was purified by silica gel chromatography (CH₂Cl₂/CH₃OH, 20:1) to give compound **12a** (1.10 g, 80 %) as a brown solid. MS (ESI) m/z: [M+H]⁺

=461.2.

4.1.3 (S)-tert-Butyl 3-((6-(6-methoxy-5-nitropyridin-3-yl)quinazolin-4-yl)amino) piperidine-1-carboxylate (12b)

To a solution mixture of **11** (500 mg, 1.23mmol) in DME/H₂O (8 mL/2 mL), were added 2-methoxy-3-nitro-5-(4,4,5,5-tetramethyl-1,3,2-dioxaborolan-2-yl)pyridine (518.2 g, 1.85 mmol), PdCl₂(dppf) (75 g, 0.12 mmol) and Na₂CO₃ (182 mg, 1.72 mmol). The mixture was refluxed under N₂ atmosphere for 4 h and then the mixture was concentrated and purified by silica gel chromatography (CH₂Cl₂/CH₃OH, 20:1) to give compound **12b** (531 mg, 90 %) as a brown solid. MS (ESI) m/z: $[M+H]^+ = 481.2$.

4.1.4

(S)-tert-Butyl

3-((6-(6-methoxy-5-(trifluoromethyl)pyridin-3-yl)quinazolin-4-yl)amino)piperidine-1carboxylate (12c)

To a solution mixture of **11** (350 mg, 0.86 mmol) in DME/H₂O (8 mL/2 mL), were added the 2-methoxy-5-(4,4,5,5-tetramethyl-1,3,2-dioxaborolan-2-yl)-3-(trifluoromethyl) pyridine (561 mg, 1.85 mmol), PdCl₂(dppf) (73.2 mg, 0.1 mmol) and Na₂CO₃ (182 mg, 1.72 mmol). The mixture was refluxed under N₂ atmosphere for 4 h and then the mixture was concentrated and purified by silica gel chromatography (CH₂Cl₂/CH₃OH, 20:1) to give compound **12c** (510 mg, 81 %) as a brown solid. MS (ESI) m/z: $[M+H]^+ = 504.2$.

4.1.5

(S)-2-Methoxy-5-(4-((1-(tetrahydro-2H-pyran-4-carbonyl)piperidin-3-yl)amino)quinaz olin-6-yl)nicotinonitrile (A1)

To a mixture of **12a** (200 mg, 0.43 mmol) in CH₂Cl₂ (5 mL), was added TFA (1 mL). The mixture was stirred at room temperature for 2h. The mixture was concentrated, and then the residue was dissolved in CH_2Cl_2 (5 mL), were added tetrahydro-2H-pyran-4-carboxylic acid (85 mg, 0.63 mmol) and HATU (326 mg, 0.86 mmol) and DIPEA (77.4 mg, 0.6mmol). The reaction mixture was stirred at room temperature for 12h. The mixture concentrated and purified by silica gel chromatography to give compound A1 (107 mg, 41 %) as a white solid. ¹H NMR (400 MHz, DMSO- d_6) δ 8.96 (s, 1H, Ar-H), 8.73 (s, 1H, NH), 8.63 (d, J = 19.0 Hz, 1H, Ar-H), 8.50 (d, J = 7.4 Hz, 1H, Ar-H), 8.19 (s, 1H, Ar-H), 8.09 – 8.00 (m, 1H, Ar-H), 7.79 (s, 1H Ar-H), 4.30-4.25 (m, 1H, CH), 4.08 (s, 3H, OCH_3), 4.03 – 3.81 (m, 2H,

CH₂), 3.47 - 3.36 (m, 2H, CH₂), 3.22-3.15 (m, 1H, CH₂), 3.05-2.72 (m, 2H, CH₂), 2.60-2.50 (m, 1H, CH₂), 2.15-2.10 (m, 1H, CH₂), 1.90-1.78 (m, 2H, CH₂), 1.71 - 1.46 (m, 4H CH₂×2), 1.29-1.20 (m, 2H, CH₂). ¹³C NMR (100 MHz, DMSO-*d*₆) δ 176.1, 172.7, 163.3, 159.5, 155.5, 150.2, 142.2, 132.6, 131.3, 128.9, 128.7, 120.9, 115.7, 115.4, 96.2, 66.8, 66.7, 55.4, 55.2, 54.1, 49.3, 37.3, 29.0, 18.5, 17.2, 13.0. ESI-HRMS C₂₆H₂₈N₆O₃, calcd [M+H]⁺: 473.2296, found: 473.2306.

4.1.6

(S)-2-Methoxy-5-(4-((1-(1-methylpiperidine-4-carbonyl)piperidin-3-yl)amino)quinazol in-6-yl)nicotinonitrile (A2)

Using the similar procedure of A1, A2 was prepared s a white solid. Yield: 56 %; ¹H NMR (400 MHz, DMSO-*d*₆) δ 8.94 (d, *J* = 2.2 Hz, 1H, Ar-H), 8.70 (d, *J* = 2.3 Hz, 1H, NH), 8.61 (d, *J* = 14.0 Hz, 1H, Ar-H), 8.47 (d, *J* = 4.4 Hz, 1H, Ar-H), 8.15 (d, *J* = 8.6 Hz, 1H, Ar-H), 8.04 (dd, *J* = 18.6, 6.8 Hz, 1H, Ar-H), 7.76 (d, *J* = 8.6 Hz, 1H, Ar-H), 4.30 – 4.23 (m, 1H, CH), 4.06 (s, 3H, OCH₃), 3.18 – 2.92 (m, 4H CH₂×2), 2.80 – 2.53 (m, 4H CH₂×2), 2.16 (s, 3H CH₃), 2.13 – 1.85 (m, 3H CH₂+CH), 1.84 – 1.70 (m, 2H CH₂), 1.67 – 1.58 (m, 2H CH₂), 1.52 – 1.31 (m, 2H CH₂). ¹³C NMR (100 MHz, DMSO-*d*₆) δ 171.8, 163.4, 159.5, 155.8, 155.7, 150.3, 149.4, 142.3, 132.7, 131.4, 128.9, 121.0, 115.7, 115.5, 96.3, 55.2, 53.6, 53.4, 49.3, 47.4, 46.1, 43.5, 42.1, 35.4, 30.2, 26.4, 24.4. ESI-HRMS C₂₇H₃₁N₇O₂, calcd [M+H]⁺: 486.2612, found: 486.2617.

4.1.7 (S)-2-Methoxy-5-(4-((1-(piperidine-4-carbonyl)piperidin-3-yl)amino) quinazolin-6-yl)nicotinonitrile (A3)

Using the similar procedure of **A1**, **A3** was prepared as a white solid. Yield: 56 %; ¹H NMR (400 MHz, DMSO- d_6) δ 8.99 (t, J = 1.8 Hz, 1H, Ar-H), 8.85 – 8.80 (m, 1H, NH), 8.76 (d, J = 1.8 Hz, 1H, Ar-H), 8.65 – 8.60 (m, 1H, Ar-H), 8.39 – 8.34 (m, 2H, Ar-H), 7.88 (dd, J = 5.7, 2.6 Hz, 1H, Ar-H), 4.40 – 4.35 (m, 1H, CH), 4.26 – 4.20 (m, 1H, NH), 4.06 (s, 3H, OCH₃), 4.05 – 3.95 (m, 2H, CH₂), 3.40 – 3.20 (m, 2H, CH₂), 3.18 – 2.92 (m, 4H, CH₂×2), 2.80 – 2.53 (m, 4H, CH₂), 2.20 – 2.10 (m, 1H CH), 1.95 – 1.65 (m, 6H CH₂×3), 1.60 – 1.40 (m, 1H, CH₂). ¹³C NMR (101 MHz, DMSO- d_6) δ 171.9, 163.3, 159.5, 158.6, 155.7, 150.2, 142.2, 132.6, 131.3, 128.9, 128.7, 121.1, 119.2, 115.7, 115.5, 96.2, 55.2, 49.3, 46.0, 43.1, 42.9, 42.6, 35.7, 34.6, 25.8, 25.5, 16.0. ESI-HRMS C₂₆H₂₉N₇O₂, calcd [M+H]⁺: 472.2456, found: 472.2461.

4.1.8 (S)-5-(4-((1-Alanylpiperidin-3-yl)amino)quinazolin-6-yl)-2-

methoxynicotinonitrile (A4)

Using the similar procedure of **A1**, **A4** was prepared as a white solid. Yield: 64 %; ¹H NMR (400 MHz, DMSO-*d*₆) δ 8.96 (s, 1H, Ar-H), 8.86 (d, *J* = 6.4 Hz, 1H, Ar-H), 8.73 (d, *J* = 2.0 Hz, 1H, Ar-H), 8.38 (d, *J* = 8.7 Hz, 1H, Ar-H), 8.17 (s, 2H, NH₂⁺), 7.92 (dd, *J* = 8.6, 4.8 Hz, 1H, Ar-H), 4.67 – 4.38 (m, 3H, CH+NH₂), 4.08 (s, 3H OCH₃), 4.04 –3.89 (m, 1H, CH), 3.20-2.72 (m, 2H, CH₂), 2.20-2.00 (m, 1H, CH₂), 2.00-1.80 (m, 2H, CH₂), 1.55 – 1.45 (m, 1H CH₂), 1.40-1.30 (m, 3H, CH₃), 1.30-1.25 (m, 2H, CH₂). ¹³C NMR (100 MHz, DMSO-*d*₆) δ 173.8, 163.7, 159.4, 155.5, 155.3, 149.0, 145.6, 138.2, 134.8, 131.4, 128.8, 128.7, 120.4, 115.5, 111.1, 53.8, 49.3, 48.8, 46.5, 42.4, 30.3, 24.3, 21.1. ESI-HRMS C₂₃H₂₅N₇O₂, calcd [M+H]⁺: 432.2143, found: 432.2140.

4.1.9

(S)-5-(4-((1-(Dimethylglycyl)piperidin-3-yl)amino)quinazolin-6-yl)-2-methoxynicotinon itrile (A5)

Using the similar procedure of **A1**, **A5** was prepared as a white solid. Yield: 64 %; ¹H NMR (400 MHz, DMSO-*d*₆) δ 8.94 (d, *J* = 2.2 Hz, 1H Ar-H), 8.70 (s, 1H NH), 8.62 (d, *J* = 10.4 Hz, 1H Ar-H), 8.56 – 8.47 (m, 1H Ar-H), 8.18 – 8.13 (m, 1H Ar-H), 8.04 (d, *J* = 6.6 Hz, 1H Ar-H), 7.76 (dd, *J* = 8.6, 5.4 Hz, 1H Ar-H), 4.25 – -4.10 (m, 1H, CH), 4.06 (s, 3H, OCH₃), 4.00 – 3.50 (m, 2H, CH₂), 3.45 – 3.20 (m, 2H, CH₂), 3.10 – 2.70 (m, 2H, CH₂), 2.45 (s, 3H, CH₃), 2.33 (s, 3H, CH₃), 2.11 – 1.75 (m, 2H, CH₂), 1.74 – 1.43 (m, 2H, CH₂). ¹³C NMR (100 MHz, DMSO-*d*₆) δ 163.3, 159.4, 159.4, 155.8, 150.3, 149.3, 142.3, 132.6, 131.4, 129.0, 128.8, 121.0, 115.7, 115.4, 96.2, 55.4, 55.2, 49.1, 48.5, 47.35, 46.0, 44.9, 44.7, 24.7, 24.1. ESI-HRMS C₂₄H₂₇N₇O₂, calcd [M+H]⁺: 446.2299, found: 446.2301.

4.1.10

(S)-5-(4-((1-(2-Chloroacetyl)piperidin-3-yl)amino)quinazolin-6-yl)-2-methoxynicotino nitrile (A6)

Using the similar procedure of A1, A6 was prepared as a white solid. Yield: 70 %; ¹H NMR (400 MHz, DMSO- d_6) δ 8.94 (s, 1H Ar-H), 8.71 (d, J = 2.3 Hz, 1H, NH), 8.62 (d, J = 11.2 Hz, 1H, Ar-H), 8.51 (d, J = 11.9 Hz, 1H, Ar-H), 8.16 (d, J = 8.4 Hz, 1H, Ar-H), 8.07 (dd, J = 11.6, 7.4 Hz, 1H Ar-H), 7.79 – 7.74 (m, 1H Ar-H), 4.55 – 4.50 (m, 1H, CH), 4.44 - 4.29 (m, 2H CH₂), 4.26-4.18 (m, 2H, CH₂), 4.07 (s, 3H OCH₃), 3.17 - 3.06 (m, 1H, CH₂), 2.82 - 2.65 (m, 1H, CH₂), 2.15 - 1.75 (s, 2H, CH₂), 1.78 - 1.51 (m, 2H CH₂).¹³C NMR (100 MHz, DMSO- d_6) δ 165.2, 163.3, 159.4, 155.8, 150.3, 149.2, 142.3, 132.7, 131.3, 129.0, 129.0, 128.7, 121.0, 115.7, 115.4, 96.2, 55.4, 55.2, 49.1, 47.4, 42.7, 42.5, 30.3, 24.0. ESI-HRMS C₂₂H₂₁ClN₆O₂, calcd [M+H]⁺: 437.1487, found: 437.1487.

4.1.11

(S)-5-(4-((1-(2-Bromopropionyl)piperidin-3-yl)amino)quinazolin-6-yl)-2-methoxynico tinonitrile (A7)

Using the similar procedure of **A1**, **A7** was prepared as a white solid. Yield: 35 %; ¹H NMR (400 MHz, DMSO- d_6) δ 8.93 (d, J = 2.4 Hz, 1H, Ar-H), 8.69 (d, J = 1.8 Hz, 1H, Ar-H), 8.59 (s, 1H, NH), 8.51 (d, J = 12.8 Hz, 1H, Ar-H), 8.15 (dd, J = 8.7, 1.3 Hz, 1H Ar-H), 8.05 – 8.01 (m, 1H, Ar-H), 7.76 (d, J = 8.7 Hz, 1H Ar-H), 6.89 – 6.74 (m, 1H, CH₂), 6.15 – 6.08 (m, 1H, CH₂), 5.74 – 5.60 (m, 1H, CH₂), 4.67 – 4.20 (m, 3H CH + CH₂), 4.06 (s, 3H OCH₃), 3.17 – 2.68 (m, 2H, CH₂), 2.14 – 1.85 (m, 2H, CH₂), 1.85 – 1.53 (m, 2H, CH₂), 1.28-1.07 (m, 1H, CH₂). ¹³C NMR (100 MHz, DMSO- d_6) δ 164.9, 163.3, 155.8, 150.2, 149.2, 142.2, 132.6, 131.3, 129.0, 128.7, 127.7, 120.9, 115.7, 115.4, 96.20, 55.2, 49.7, 48.7, 47.5, 30.3, 25.4, 24.1. ESI-HRMS C₂₃H₂₄BrN₆O₂, calcd [M+H]⁺: 495.1139, found: 495.1158.

4.1.12 (S)-2-Methoxy-5-(4-((1-nicotinoylpiperidin-3-yl)amino)quinazolin-6-yl) nicotinonitrile (A8)

Using the similar procedure of **A1**, **A8** was prepared as a white solid. Yield: 67 %; ¹H-NMR (400 MHz, DMSO- d_6) δ 9.30 – 8.97 (m, 2H, Ar-H), 8.74 – 8.48 (m, 5H, NH+Ar-H), 8.29 – 8.15 (m, 1H Ar-H), 7.90 – 7.76 (m, 2H, Ar-H), 7.52 – 7.40 (m, 1H Ar-H), 4.30 – 4.25 (m, 1H, CH), 4.04 (s, 3H OCH₃), 3.90 – 3.51 (m, 2H CH₂), 3.25 – 2.90 (m, 2H, CH₂), 2.23 – 2.00 (m, 1H, CH₂), 2.00 – 1.75 (m, 2H, CH₂), 1.75 – 1.60 (m, 1H, CH₂). ¹³C NMR (100 MHz, DMSO- d_6) δ 167.5, 163.4, 159.7 154.1, 154.0, 150.8, 150.7, 150.7, 150.3, 148.1, 147.9, 142.3, 135.2, 132.5, 132.2, 128.5, 126.2, 123.0, 121.7, 121.5, 115.6, 114.8, 96.2, 55.2, 53.8, 49.0, 42.1, 18.5, 17.2. ESI-HRMS C₂₆H₂₃N₇O₂, calcd [M+H]⁺: 466.1986, found: 466.1984.

4.1.13 (S)-2-Methoxy-5-(4-((1-(5-nitrofuran-2-carbonyl)piperidin-3-yl)amino)

quinazolin-6-yl)nicotinonitrile (A9)

Using the similar procedure of **A1**, **A9** was prepared as a white solid. Yield: 36 %; ¹H NMR (400 MHz, DMSO-*d*₆) δ 8.93 (s, 1H, Ar-H), 8.70 (s, 1H, Ar-H), 8.60 (s, 1H, Ar-H), 8.50 (s, 1H, NH), 8.15 (d, *J* = 8.4 Hz, 1H, Ar-H), 8.01 (s, 1H, Ar-H), 7.79 (d, *J* = 3.8 Hz, 1H, Ar-H), 7.75 (d, *J* = 8.0 Hz, 1H, Ar-H), 7.58 (s, 1H Ar-H), 4.38 – 4.30 (m, 1H, CH), 4.29 – 4.09 (m, 2H, CH₂), 4.06 (s, 3H, OCH₃), 3.40 – 3.30 (m, 2H CH₂), 2.20 – 2.13 (m, 1H, CH₂), 1.98 – 1.75 (m, 2H, CH₂), 1.70 – 1.65 (m, 1H, CH₂). ¹³C NMR (100 MHz, DMSO-*d*₆) δ 163.3, 159.3, 157.4, 155.4, 151.6, 150.3, 149.3, 142.3, 140.8, 132.7, 131.3, 129.0, 128.8, 121.0, 116.9, 115.7, 115.3, 113.3, 96.2, 55.4, 55.2, 50.6, 43.5, 29.8, 23.8. ESI-HRMS C₂₅H₂₁N₇O₅, calcd [M+H]⁺: 500.1677, found: 500.1679.

4.1.14 (S)-2-Methoxy-5-(4-((1-(2,2,2-trifluoroacetyl)piperidin-3-yl)amino) quinazolin-6-yl)nicotinonitrile (A10)

Using the similar procedure of **A1**, **A10** was prepared as a white solid. Yield: 20 %; ¹H NMR (400 MHz, DMSO-*d*₆) δ 8.99 (d, *J* = 1.5 Hz, 1H, Ar-H), 8.83 (d, *J* = 2.3 Hz, 1H, Ar-H), 8.65 (dd, *J* = 6.4, 1.6 Hz, 1H, Ar-H), 8.52 (d, *J* = 4.0 Hz, 1H, NH), 8.20 (dd, *J* = 8.7, 1.8 Hz, 1H, Ar-H), 8.19 – 8.10 (m, 1H, Ar-H), 7.80 (d, *J* = 8.7 Hz, 1H, Ar-H), 4.38 – 4.30 (m, 1H, CH), 4.11 (s, 3H OCH₃), 3.33 – 3.10 (m, 2H CH₂), 2.15 – 2.05 (m, 1H, CH₂), 2.00 – 1.94 (m, 1H, CH₂), 1.88 – 1.75 (m, 1H, CH₂), 1.70 – 1.55 (m, 1H, CH₂), 1.30 – 1.20 (m, 2H, CH₂). ¹³C NMR (100 MHz, DMSO-*d*₆) δ 163.4, 159.6, 159.5, 155.4, 150.3, 142.4, 133.0, 133.0, 131.7, 129.0, 128.2, 128.1, 121.1, 115.7, 115.2, 96.3, 55.2, 54.1, 42.3, 18.5, 17.2, 13.0. ESI-HRMS C₂₂H₁₉F₃N₆O₂, calcd [M+H]⁺: 457.1594, found: 457.1597.

4.1.15 Ethyl (S)-2-(3-((6-(5-cyano-6-methoxypyridin-3-yl)quinazolin-4-yl)amino) piperidin-1-yl)acetate (A11)

Using the similar procedure of A1, A11 was prepared as a white solid. Yield: 57 %; ¹H NMR (400 MHz, DMSO- d_6) δ 8.92 (d, J = 2.5 Hz, 1H), 8.69 (d, J = 2.5 Hz, 1H, Ar-H), 8.58 (d, J = 1.6 Hz, 1H, Ar-H), 8.47 (s, 1H, NH), 8.13 (dd, J = 8.7, 1.8 Hz, 1H, Ar-H), 7.91 (d, J = 7.8 Hz, 1H, Ar-H), 7.73 (d, J = 8.7 Hz, 1H, Ar-H), 4.43 – 4.33 (m, 1H, CH), 4.12 – 4.06 (m, 2H CH₂), 4.06 (s, 3H, OCH₃), 3.32 – 3.25 (m, 2H, CH₂), 3.08 – 3.00 (m, 1H, CH₂), 2.81 – 2.75 (m, 1H, CH₂), 2.36 – 2.30 (m, 2H, CH₂), 2.01 – 1.91 (m, 1H, CH₂), 1.79 – 1.71 (m, 1H, CH₂), 1.64 – 1.38 (m, 2H CH₂), 1.18 (t, J = 7.1 Hz, 3H, CH₃). ¹³C NMR (100 MHz, DMSO-*d*₆) δ 170.7, 163.3, 159.3, 155.9, 150.2, 149.2, 142.2, 132.5, 131.1, 129.0, 128.7, 120.8, 115.7, 115.4, 96.2, 60.3, 58.9, 57.3, 55.2, 52.4, 47.9, 29.9, 24.5, 14.6. ESI-HRMS C₂₄H₂₆N₆O₃, calcd [M+H]⁺: 447.2139, found: 447.2132.

4.1.16 Ethyl (S)-2,2,2-trifluoro-1-(3-((6-(6-methoxy-5-nitropyridin-3-yl)quinazolin-4-yl) amino)piperidin-1-yl)ethan-1-one (A12)

Using the similar procedure of **A1**, **A12** was prepared as a white solid. Yield: 52 %; ¹H NMR (400 MHz, DMSO-*d*₆) δ 8.99 (d, *J* = 1.5 Hz, 1H, Ar-H), 8.82 (d, *J* = 2.3 Hz, 1H, NH), 8.65 (dd, *J* = 6.4, 1.6 Hz, 1H, Ar-H), 8.52 (d, *J* = 4.0 Hz, 1H, Ar-H), 8.20 (dd, *J* = 8.7, 1.8 Hz, 1H, Ar-H), 8.15 – 8.06 (m, 1H, Ar-H), 7.79 (d, *J* = 8.7 Hz, 1H, Ar-H), 4.43 – 4.33 (m, 1H, CH), 4.10 (s, 3H OCH₃), 3.33 – 3.11 (m, 2H CH₂), 2.14 – 2.00 (m, 1H, CH₂), 2.00 – 1.93 (m, 1H, CH₂), 1.90 – 1.75 (m, 1H, CH₂), 1.65 – 1.50 (m, 1H, CH₂), 1.27 – 1.20 (m, 2H, CH₂). ¹³C NMR (100 MHz, DMSO-*d*₆) δ 159.5, 159.4, 155.8, 155.4, 150.2, 149.3, 134.3, 133.4, 132.4, 131.5, 129.3, 128.8, 121.2, 121.2, 115.4, 55.3, 47.9, 47.2, 43.8, 24.8, 23.4. ESI-HRMS C₂₁H₁₉F₃N₆O₄, calcd [M+H]⁺: 477.1493, found: 477.1493.

4.1.17

Ethyl

(S)-2-(3-((6-(5-cyano-6-methoxypyridin-3-yl)quinazolin-4-yl)amino)piperidin-1-yl)acet ate (A13)

Using the similar procedure of **A1**, **A13** was prepared as a white solid. Yield: 50 %; ¹H NMR (400 MHz, DMSO- d_6) δ 8.92 (s, 1H Ar-H), 8.62 (s, 1H NH), 8.49 (s, 1H Ar-H), 8.45 (s, 1H Ar-H), 8.17 (d, J = 8.5 Hz, 1H Ar-H), 7.96 (d, J = 7.6 Hz, 1H Ar-H), 7.76 (d, J = 8.6 Hz, 1H Ar-H), 4.43 – 4.33 (m, 1H, CH), 4.12 – 4.06 (m, 2H CH₂), 4.06 (s, 3H, OCH₃), 3.32 – 3.25 (m, 2H, CH₂), 3.08 – 3.00 (m, 1H, CH₂), 2.81 – 2.75 (m, 1H, CH₂), 2.36 – 2.30 (m, 2H, CH₂), 2.01 – 1.91 (m, 1H, CH₂), 1.79 – 1.71 (m, 1H, CH₂), 1.64 – 1.38 (m, 2H CH₂), 1.19 (t, J = 7.1 Hz, 3H, CH₃). ¹³C NMR (100 MHz, DMSO- d_6) δ 170.7, 160.0, 159.3, 155.8, 149.8, 149.2, 135.7, 133.1, 131.5, 128.8, 128.7, 122.2, 121.0, 115.5, 111.8, 60.3, 58.9, 57.3, 54.8, 52.4, 47.9, 29.9, 24.5, 14.6. ESI-HRMS C₂₄H₂₆F₃N₅O₃, calcd [M+H]⁺: 490.2061, found: 490.2058.

4.1.18

(S)-5-(4-((1-(2-Hydroxyethyl)piperidin-3-yl)amino)quinazolin-6-yl)-2-methoxynicotino nitrile (A14) Using the similar procedure of **A1**, **A14** was prepared as a white solid. Yield: 53 %; ¹H NMR (400 MHz, DMSO-*d*₆) δ 8.95 (s, 1H Ar-H), 8.74 – 8.68 (m, 1H Ar-H), 8.64 – 8.58 (m, 1H Ar-H), 8.51 – 8.47 (m, 1H Ar-H), 8.19 – 8.13 (m, 1H Ar-H), 8.07 – 7.95 (m, 1H NH), 7.78 – 7.73 (m, 1H Ar-H), 4.55 – 4.35 (m, 1H CH), 4.07 (s, 3H OCH₃), 3.60 – 3.50 (m, 2H CH₂), 3.40 – 3.30 (m, 2H CH₂), 3.07 (s, 1H OH), 2.75 – 2.65 (m, 2H CH₂), 2.40 – 2.25 (m, 2H, CH₂), 2.05 – 1.90 (m, 1H, CH₂), 1.85 – 1.75 (m, 1H, CH₂), 1.70 – 1.45 (m, 1H, CH₂). ¹³C NMR (100 MHz, DMSO-*d*₆) δ 163.5, 160.1, 153.8, 150.4, 142.4, 134.1, 133.1, 128.5, 121.6, 119.0, 116.0, 115.6, 114.6, 113.1, 96.3, 58.9, 55.4, 55.3, 52.3, 46.08, 39.9, 27.8, 21.6. ESI-HRMS C₂₂H₂₄N₆O₂, calcd [M+H]⁺: 405.2034, found: 405.2062.

4.2 PI3K enzymatic activity assay

The PI3K isoform activity assay was performed as our previous procedures. ¹⁸ All the PI3K isoform reaction was conducted at 30 °C for 60 min. The 50 μ L reaction mixture contains 50 mM HEPES (pH 7.5), 50 mM NaCl, 3 mM MgCl₂, 0.025 mg/mL BSA, 0.2 μ g/mL PI3K, 10 μ M ATP and 100 μ M PI substrate. The tested compounds were diluted in 10% DMSO and 5 μ L of the dilution was added to a 50 μ L reaction so that the final concentration of DMSO is 1% in all the reactions. After the plate was cooled at room temperature for 5 min, 5 μ L of ADP-Glo reagent was added into each well to stop the reaction and consume the remaining ADP within 40 min. At the end, 10 μ L of kinase detection reagent was added into the well and incubated for 30 min. The assay was performed using Kinase-Glo Plus luminescence kinase assay. The luminescence signal was detected by VICTOR-X multi-label plate reader. The IC₅₀ values were calculated using nonlinear regression with normalized dose–response fit using GraphPad Prism 5.0 sofeware.

4.3 Anti-proliferative assays using a MTT method in vitro

Cellular proliferation was determined using a MTT method in vitro.¹⁹

4.4 Western blot assay

The levels of Akt and phospho-Akt (p-Akt, S473) in SU-DHL-6 cells were determined by Western blot. SU-DHL-6 cells were treated with final concentrations of 1000 nM, 100 nM, and 10 nM of **A8** and with the concentrations of 1000 and 100 nM of idelalisib for 8 h. Cell lysates were clarified by centrifugation at 12,000 rpm for 20 min at

4 °C, and supernatant was collected. Equal amounts of protein were subjected to SDS-PAGE and transferred to nitrocellulose membranes (Merck Millipore, Billerica, MA, USA). The membranes were blocked and incubated overnight at 4 °C with primary antibodies against p-Akt, Akt, and beta-actin, followed by incubation with appropriate secondary antibodies. Antibody binding was detected with chemiluminescence reagents (Sigma-Aldrich, St. Louis, MO, USA).

4.5 Molecular docking simulation

The molecular docking was performed as our previous reported methods.¹⁹

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Appendix A. Supplementary data

Supplementary data to this article can be found attached.

References

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Synthesis and Biological Evaluation of 4-(Piperid-3-yl)amino Substituted 6-Pyridylquinazolines as Potent PI3Kδ Inhibitors

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Graphical Abstract:

IC₅₀ = 0.7 nM (PI3Kdelta)

IC₅₀ = 0.12 uM (SU-DHL-6)



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