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Design and Synthesis of Potent Quinazolines as Selective β -Glucocerebrosidase Modulators

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Abstract

Gaucher's disease is a common genetic disease caused by mutations in the β -glucocerebrosidase (GBA1) gene that have been also linked to increased risk of Parkinson's disease and Lewy Body Dementia. Stabilization of misfolded mutant β -glucocerebrosidase (GCase) represents an important therapeutic strategy in synucleinopathies. Here we report a novel class of GCase quinazoline inhibitors, obtained in a high throughput screening, with moderate potency against wild-type GCase. Rational design and a SAR study of this class of compounds led to a new series of quinazoline derivatives with single digit nanomolar potency. These compounds were shown to selectively stabilize GCase when compared to other lysosomal enzymes and to increase N370S mutant GCase protein concentration and activity in cell assays. To the best of our knowledge, these molecules are the most potent non-iminosugar GCase inhibitors to date that may prove useful for future mechanistic studies and therapeutic approaches in Gaucher's and Parkinson's diseases.

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The authors declare no competing financial interest.

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Supporting Information Available: ¹H and ¹³C NMR spectra of **4**, **6c**, **7a**, **7b**, **8c**, **9a**, **9b**, **9c**, **11b**, **11d**, **11f**, **11g**, and **11h** (PDF) Molecular formula strings (CSV)

Keywords

β-Glucocerebrosidase; Gaucher's disease; Parkinson's disease; Quinazoline; Modulator; SAR

INTRODUCTION

Gaucher's disease (GD), the most common lysosomal storage disease, is caused by a recessively inherited deficiency in β -glucocerebrosidase (GCase) and subsequent accumulation of glucoceramides, toxic lipid substrates^{1,2}. Substrate accumulation leads to hepatosplenomegaly, bone marrow suppression, and bone lesions^{1–3}. Many of the GCase mutations are missense mutations⁴ that result in single amino acid substitutions of the enzyme. Most of these mutations, including the prevalent N370S mutation, are still functional, although with very low residual GCase activity⁵ due to enzyme misfolding, and proteasome-mediated breakdown⁵. Current treatments for GD include enzyme replacement therapy (ERT) and substrate reduction therapy (SRT) ^{2,6}. In recent years, mutations in GBA1 were also found to be a major risk factor for Parkinson's disease (PD) and dementia with Lewy bodies (DLB)^{7–11}. Accumulation of β -glucosylceramide, the substrate of GCase, in neurons promotes the formation of α -synuclein oligomers, which are considered toxic in PD¹². Enhancement of GCase activity is thought to be a potential therapeutic strategy for GCase-associated synucleinopathies, including PD^{13,14}.

An emerging therapeutic approach involves the restoration of proper folding and lysosome delivery of degradation-prone mutant enzymes using small molecules as pharmacological chaperones (PCs)⁵. Previous studies have shown that iminosugars increase the cellular activity of the N370S mutant form of GCase^{15,16}, as well as of the wild-type enzyme^{5,17}. Isofagomine (IFG, **1**) attracted the most attention in the iminosugar class of compounds (Figure 1)¹⁸. However, iminosugars tend to have poor selectivity and relatively short half-lives in cells¹⁹. Several different scaffolds of non-iminosugar inhibitors (**2** and **3** are examples in Figure 1) have been reported as GCase PCs since 2007^{20–24}; however, the binding site and the interaction with GCase of these non-iminosugar PCs remain unknown.

In our high throughput screening effort to discover potent GCase modulators, compound 4 (Figure 1) was identified as a potent GCase inhibitor (IC₅₀ 0.177 μ M) in a 4methylumbelliferyl β -D-glucopyranoside (4MU- β -Glc) enzyme activity based high throughput screen. The activity was confirmed with additional synthesized compounds. To further develop potent GCase modulators and use them to explore the properties of the binding site, we carried out a structure activity relationship (SAR) study of a series of quinazoline derivatives, leading to the discovery of single digit nanomolar potency GCase inhibitory modulators.

CHEMISTRY

The synthesis of compound **4** and its analogues for SAR exploration is straightforward and is detailed in Scheme 1 and 2. As showed in Scheme 1, **5** was prepared from 2-aminobenzonitrile and nicotinoyl chloride according to a known method²⁵. The reaction of **5** and

appropriate amines in the presence of potassium carbonate as a base afforded **4**, **6a-6i**, **7a-7i**, **8a-8g**, and **9a-9f**.

Additional analogues having modifications at position 2 of the quinazoline ring were synthesized, employing alkylation of 2,3-dihydro-1*H*-inden-2-amine with 2,4-dichloroquinazoline, followed by Suzuki coupling with appropriate boronic acids to afford **11a-11h** (Scheme 2). The structure and purity of all the prepared compounds were confirmed by spectroscopic and analytical techniques.

RESULTS AND DISCUSSION

In our high throughput screening efforts to discover potent GCase inhibitors/activators, we used recombinant wild-type GCase and 4MU- β -Glc as substrate in an optimized pH 5.9 buffer²⁶. In previous studies, high concentrations of taurocholate (4–10 mM) were used to improve the signal in GCase enzyme activity assays^{16,27,28}. We found that taurocholate can interfere with our assay results, which was also reported recently by Berger et al.²⁹; therefore, we excluded taurocholate in our screening assay. Using this approach, we discovered several different scaffolds of GCase inhibitors and activators with moderate activity. Among these, we identified a quinazoline compound (**4**, Figure 1) as a potent GCase inhibitor. The quinazoline ring had previously been found as the best scaffold for GCase inhibitors among several ring systems assayed²¹. Here we describe our modifications of the substituents on the quinazoline ring.

To examine the SAR at the amino group, a series of substituents was introduced at the 4position of the *N*-cyclohexyl ring of **4** (Table 1). A 4-methyl substituent (**6a**, *cis/trans* = 3/2mixture) resulted in 3-fold higher activity (IC₅₀ 56 nM), while 4-ethyl substitution (**6b**, *cis/ trans* = 3/2 mixture) did not show a significant change of activity, suggesting that a smaller hydrophobic group at this position may be beneficial. Then, a *trans*-4-methyl compound (**6c**) was synthesized, which exhibited great improvement of inhibitory activity (IC₅₀ 20 nM). Further modification by installation of 4-methoxyl (**6d**), 4,4-dimethyl (**6e**), or 4,4-difluoro (**6f**) groups at the same position decreased the potency; amino substituents were detrimental to activity. Replacement of the cyclohexyl group by tetrahydro-2*H*-pyran-3-yl (**6h**), or *N*substituted piperidine (**6i-6k**) afforded weak or inactive compounds, supporting the importance of the lipophilic cyclohexyl ring.

To further expand the SAR of the amino group, the cyclohexyl ring of **4** was replaced by a series of saturated carbon rings of different sizes. A dramatic SAR was observed with different carbon rings (**7a-7e**). As shown in Table 2, the larger cycloalkyl rings were more potent; the compound with a cyclooctyl group (**7a**, IC₅₀ 27 nM) was the most potent. However, when bulk was introduced to the cycloalkyl ring, the potency of the compounds (**7f** and **7g**) decreased, suggesting that the hydrophobic binding pocket may be compact. Introduction of one or two carbons between the cyclohexyl and NH groups (**7h** and **7i**) in **4** decreased the inhibitory activity, again indicating a hydrophobic pocket with limited volume.

To understand the nature of the binding site, a phenyl ring with different length linkers was introduced into the molecules (**8a-e**, Table 3). Compound **8a**, by replacement of the

cyclohexyl ring in **4** with a phenyl ring, lost activity. Insertion of a 1–4 carbon linker between the phenyl and quinazoline rings gave **8b-8e**. Interestingly, **8c**, with a phenylethyl group, was slightly more potent than **4**. Extension of the linker did not benefit activity, suggesting that a two carbon length linker between the phenyl ring and secondary amino group of **8c** may allow optimal binding of the phenyl group. Substitution of the phenyl group in **8** with a 2- or 3-pyridine ring (**8f** or **8g**) sharply diminished potency, indicating a repulsive effect of the pyridine nitrogen atom.

To enhance the binding affinity of the compounds with GCase, a new series of compounds was designed to integrate both the hydrophobic interaction of the A ring and the π - π interaction of the B ring by fusing the cyclohexyl ring (4) with a phenyl ring (Figure 2 and Table 4). These derivatives (**9a-c**) exhibited single digit nanomolar inhibitory activity against GCase. Stereochemistry did not seem to be important (**9a** and **9b**). The compound with an indane ring (**9c**) gave comparable activity to that of the tetralin ring (**9a**). Attachment of the quinazoline ring to the tetralin ring (**9d**) and indane ring (**9e**) at position 1 instead of position 2 (**9a** and **9c**) dramatically increased the IC₅₀ values to the low micromolar range, indicating the importance of the orientation of this substituent for binding activity of these inhibitors. The introduction of an oxygen atom to give a chromane (**9f**) did not significantly affect the potency. These results suggest important hydrophobic and π - π interactions in the binding of this series of compounds to GCase.

Finally, we examined the substituent effect on the pyridine ring. A methyl group was introduced at different positions of the 3-pyridinyl ring of **9c** to give **11a-c** (Table 5). Methylation of the pyridine ring decreased the potency. Whereas the compound with a methyl group at position 4 of the pyridinyl ring (**11a**) showed only moderate potency, the other two compounds (**11b** and **11c**) with a methyl group at positions 5- and 6 were much more potent, comparable to that when a 2-furanyl group replaced the pyridine ring, but still not as potent as **9c**. The replacement of the 3-pyridinyl ring in **9c** with either phenyl or 3-thienyl groups, however, gave compounds that retained the same potency as **9c**, suggesting that more groups could be introduced at position 2 of the quinazoline ring.

We also evaluated the activity of these selected compounds at various pH values (Table 6). Interestingly, compared to the inhibitory activity at pH 5.9, the activity of **9a**, **9b**, and **9c** with a 3-pyridinyl ring decreased by 3 fold at pH 5.0, while the activity of **11d**, **11f**, and **11g** only dropped slightly at both pH conditions. Although the cause for the change in IC_{50} values is not clear, an explanation may involve the protonation state of the compounds or the variability of the enzyme efficacy at different pH values.

Compounds shown to act as pharmacological chaperones for GCase (or other lysosomal enzymes) also stabilize the enzyme against thermal denaturation. A fluorescent thermal shift assay was developed to evaluate the binding affinity of ligands with protein³⁰. To evaluate their abilities to stabilize GCase, the most potent compounds, **9a**, **9b**, **11d**, **11g** and **11f** were accessed in a wild-type GCase fluorescent thermal shift assay with a negative control (**8a**) and a positive control (IFG) at pH 5.0. The selected compounds increased the GCase melting point in a dose-dependent manner (Figure 3), while inactive compound **8a** did not change the melting points significantly. Most compounds exhibited greater ability to stabilize GCase

than IFG at lower concentrations, and **11g** showed the maximum thermal shift up to around 11 °C. It should be noted that the increment in melting point has a direct correlation with the compound's binding affinity³¹. The maximum thermal shift of our compounds corresponded to their inhibitory activity at pH 5.0, suggesting a close correlation of the binding affinity and the inhibitory activity for these compounds.

The selected compounds (9a, 9b, 11d, 11g, and 11f) were further evaluated against two other lysosomal hydrolases, acid α -glucosidase (GAA) and α -galactosidase A (GLA). The activities of tested enzymes were not significantly changed by compound treatments up to 100 μ M (representative results shown for 11g in Figure 4).

We further tested compounds **4** and **11g** by measuring GCase activity at various substrate concentrations (30–150 μ M) and in the absence or presence of increasing concentrations of GCase inhibitors. Similarly to reported non-iminosugar inhibitors²⁰, both of our inhibitors exhibited linear mixed inhibition, with an increase in K_m and decrease in V_{max} values upon increasing inhibitor concentrations (Figure 5A and 5B)

Finally, we tested **11g** in patient-derived N370S fibroblasts. The fibroblasts were treated with **11g** and IFG for 3 days, and levels of GCase protein were determined by Western blot. We found that **11g** treatments (2, 5, and 10 μ M) significantly increased the concentration of GCase, while IFG showed a similar effect only at the higher concentrations (20 and 50 μ M) (Figure 6, top). We also used endoH and PNGase F to digest the cell lysates to determine whether **11g** treatment affected ER maturation. The Western blot (Figure 6, middle) showed that most of the GCase bands were resistant to endoH digestion, indicating that the major GCase signal after treatment is a post-ER form. PNGase F digestion gave a single band with increased GCase concentration after **11g** treatment (Figure 6, bottom). In addition to this finding, we also found that a 3-day treatment of compound **11g** (2 μ M) increased 50% of GCase levels and the activity of patient-derived N370S fibroblasts.

CONCLUSIONS

In this paper, we describe the design and SAR of a series of quinazoline GCase inhibitors having single digit nanomolar potency. The SAR suggested that a hydrophobic interaction and a π - π interaction may be involved in compound binding to GCase. These quinazoline derivatives also stabilized GCase, as indicated by thermal shift assays, and exhibited high selectivity against other lysosomal hydrolases. Furthermore, the most potent compound (**11g**) increased the mature post-ER form of GCase and the enzyme activity in patient-derived N370S fibroblasts. It will be of interest to further test these compounds in other biological assays and models of Gaucher's and Parkinson's disease.

EXPERIMENTAL SECTION

Materials and Methods

Chemistry—Commercially available reagents and solvents were used without further purification. All reactions were monitored by thin-layer chromatography (TLC) using 0.25

mm Silicycle extra hard 250 μ M TLC plates (60 F254). Purification of reaction products was carried out by flash chromatography using an Agilent 971-FP flash purification system with Silicycle silica gel columns. The yields are not optimized. The purity of all compounds was over 95% as analyzed with an Agilent 1260 Infinity HPLC system and an Agilent Poroshell 120 EC-C18 (4.6 × 50 mm, 2.7 μ m) reverse phase column, detecting with UV absorbance (254 nm). ¹H NMR and ¹³C NMR spectra were obtained using a Bruker Avance III 500 MHz system (500 MHz for ¹H NMR and 125 MHz for ¹³C NMR) spectrometer. Chemical shifts are reported relative to chloroform (δ = 7.26 for ¹H NMR and δ = 77.16 for ¹³C NMR spectra) or dimethyl sulfoxide (δ = 2.50 for ¹H and δ = 39.52 for ¹³C NMR spectra). Data are reported as br = broad, s = singlet, d = doublet, t = triplet, q = quartet, m = multiplet. Mass spectra (HRMS) were performed using an Agilent 6210A LC-TOF instrument with a dual spray ESI source, with a high resolution Time of Flight (TOF) Mass analyzer and collecting in a 2GigHz detector mode, coupled with an Agilent 1200 HPLC.

Preparation of 4-chloro-2-(pyridin-3-yl)quinazoline (5)²⁵—To a solution of 2aminobenzonitrile (5.90 g, 50 mmol) in sulfolane (20 mL) was added nicotinoyl chloride hydrochloride (12.0 g, 67.4 mmol), and the mixture was stirred at 100°C for 16 h. PCl₅ (18.2 g, 87.5 mmol) was added in one portion and stirred at 100 °C for 10 h. The mixture was cooled to room temperature, and carefully poured into 400 mL of saturated sodium bicarbonate solution cooling in an ice bath. The solid was filtered, washed with water, dried, and purified by flash chromatography to give **5** as a pale-yellow solid (5.50 g, 46%); mp 160–163 °C. ¹H NMR (400 MHz, CDCl₃) & 9.76 (d, *J* = 1.3 Hz, 1H), 8.82 (dt, *J* = 8.0, 1.9 Hz, 1H), 8.73 (dd, *J* = 4.7, 1.4 Hz, 1H), 8.26 (dd, *J* = 8.4, 0.8 Hz, 1H), 8.10 (d, *J* = 8.4 Hz, 1H), 7.95 (ddd, *J* = 8.4, 7.0, 1.4 Hz, 1H), 7.69 (ddd, *J* = 8.2, 7.0, 1.1 Hz, 1H), 7.44 (dd, *J* = 7.5, 4.8 Hz, 1H). ¹³C NMR: (100 MHz, CDCl₃) & 162.9, 158.3, 151.8, 151.7, 150.3, 136.1, 135.2, 132.4, 129.1, 128.9, 126.0, 123.5, 122.8. ESI-MS m/z: 242 (M+H)⁺.

General Procedure for Compound 4, 6a-6i, 7a-7i, 8a-8g and 9a-9f—A mixture of 4-chloro-2-(pyridin-3-yl)quinazoline **5** (72 mg, 0.3 mmol), amine (0.3 mmol), and potassium carbonate (69 mg, 0.3 mmol) in DMF (3 mL) was stirred at room temperature or 60 °C overnight. Water (20 mL) was added, and the formed solid was filtered, washed with water, and dried *in vacuo* to give product. The products were usually pure (>95% purity). Those products without sufficient purity were purified by flash chromatography.

Analytical Data for Compounds

N-Cyclohexyl-2-(pyridin-3-yl)quinazolin-4-amine (4)—Off-white solid (64 mg, 70%); mp 108–111 °C. ¹H NMR (500 MHz, CDCl₃) δ 9.74 (s, 1H), 8.84 (d, *J* = 6.6 Hz, 1H), 8.70 (s, 1H), 7.95 (d, *J* = 7.1 Hz, 1H), 7.74 (t, *J* = 7.7 Hz, 2H), 7.49 – 7.35 (m, 2H), 5.77 (s, 1H), 4.48 – 4.32 (m, 1H), 2.28 – 2.20 (m, 2H), 1.93 – 1.79 (m, 2H), 1.79 – 1.67 (m, 1H), 1.61 – 1.46 (m, 2H), 1.46 – 1.23 (m, 3H). ¹³C NMR (125 MHz, CDCl₃) δ 158.9, 158.8, 150.7, 150.5, 150.3, 135.7, 134.6, 132.7, 129.0, 125.8, 123.2, 120.5, 113.9, 50.1, 33.0, 25.9, 25.1. HRMS (ESI): calcd for C₁₉H₂₁N₄ [M+H]⁺, 305.1761; found, 305.1764.

N-(4-Methylcyclohexyl)-2-(pyridin-3-yl)quinazolin-4-amine (*cis/trans* = 3/2) (6a) —White solid (54 mg, 43%); mp 130–134 °C.

Cis isomer: ¹H NMR (500 MHz, CDCl₃) δ 9.75 (s, 1H), 8.80 (d, *J* = 7.8 Hz, 1H), 8.69 (s, 1H), 7.92 (d, *J* = 8.3 Hz, 1H), 7.77 – 7.71 (m, 2H), 7.48 – 7.39 (m, 2H), 5.87 (d, *J* = 5.8 Hz, 1H), 4.65 – 4.58 (m, 1H), 2.02–1.94 (m, 2H), 1.86–1.81 (m, 2H), 1.76 – 1.62 (m, 3H), 1.36 – 1.21 (m, 2H), 1.00 (d, *J* = 6.5 Hz, 3H). ¹³C NMR (125 MHz, CDCl₃) δ 159.0, 158.8, 150.8, 150.4, 150.3, 135.8, 134.5, 132.8, 128.8, 125.9, 123.3, 120.5, 113.9, 47.2, 30.8, 30.4, 29.2, 21.3.

Trans isomer: ¹H NMR (500 MHz, CDCl₃) δ 9.75 (s, 1H), 8.80 (d, J = 7.8 Hz, 1H), 8.69 (s, 1H), 7.92 (d, J = 8.3 Hz, 1H), 7.77 – 7.70 (m, 2H), 7.48 – 7.39 (m, 2H), 5.63 (d, J = 5.3 Hz, 1H), 4.38 – 4.26 (m, 1H), 2.28 (d, J = 9.9 Hz, 2H), 1.86–1.81 (m, 2H), 1.54 – 1.40 (m, 1H), 1.40 – 1.30 (m, 2H), 1.30 – 1.14 (m, 2H), 0.98 (d, J = 6.5 Hz, 3H). ¹³C NMR (125 MHz, CDCl₃) δ 159.1, 158.8, 150.8, 150.5, 150.4, 135.8, 134.5, 132.8, 129.0, 125.9, 123.3, 120.7, 114.0, 50.5, 34.1, 33.1, 32.3, 22.4. HRMS (ESI): calcd for C₂₀H₂₃N₄ [M+H]⁺, 319.1917; found, 319.1921.

N-(4-Ethylcyclohexyl)-2-(pyridin-3-yl)quinazolin-4-amine (*cis/trans* = 3/2) (6b)—Pale-yellow solid (38 mg, 30%); mp 62–65 °C.

Cis isomer: ¹H NMR (500 MHz, CDCl₃) δ 9.74 (s, 1H), 8.81 (d, *J* = 7.7 Hz, 1H), 8.68 (s, 1H), 7.92 (d, *J* = 8.2 Hz, 1H), 7.77 – 7.71 (m, 1H), 7.48 – 7.39 (m, 2H), 5.85 (d, *J* = 5.8 Hz, 1H), 4.67 – 4.61 (m, 1H), 2.02–1.70 (m, 5H), 1.38 – 1.15 (m, 4H), 0.93 (t, *J* = 7.4 Hz, 3H). ¹³C NMR (125 MHz, CDCl₃) δ 159.0, 158.8, 150.8, 150.4, 135.8, 134.5, 132.8, 128.9, 125.9, 123.3, 120.5, 113.9, 47.5, 39.0, 30.8, 29.7, 29.3, 28.1, 11.8.

Trans isomer: ¹H NMR (500 MHz, CDCl₃) δ 9.74 (s, 1H), 8.81 (d, J = 7.7 Hz, 1H), 8.68 (s, 1H), 7.92 (d, J = 8.2 Hz, 1H), 7.77 – 7.71 (m, 1H), 7.48 – 7.39 (m, 2H), 5.63 (brs, 1H), 4.38 – 4.30 (m, 1H), 2.28 (d, J = 9.9 Hz, 2H), 1.99–1.70 (m, 2H), 1.54 – 1.40 (m, 1H), 1.40 – 1.30 (m, 2H), 1.30 – 1.15 (m, 2H), 0.93 (t, J = 7.4 Hz, 3H). ¹³C NMR (125 MHz, CDCl₃) δ 159.1, 158.8, 150.8, 150.4, 135.8, 134.5, 132.8, 129.0, 125.9, 123.3, 120.7, 114.0, 50.8, 37.7, 33.1, 31.7, 28.3, 11.8. HRMS (ESI): calcd for C₂₁H₂₅N₄ [M+H]⁺, 333.2074; found, 333.2080.

Trans-N-(4-Methylcyclohexyl)-2-(pyridin-3-yl)quinazolin-4-amine (6c)—White solid (52 mg, 55%); mp 173–175 °C. ¹H NMR (500 MHz, CDCl₃) δ 9.75 (s, 1H), 8.80 (d, J = 7.8 Hz, 1H), 8.69 (s, 1H), 7.92 (d, J = 8.3 Hz, 1H), 7.77 – 7.71 (m, 1H), 7.70 (d, J = 8.2 Hz, 1H), 7.53 – 7.33 (m, 2H), 5.63 (d, J = 5.3 Hz, 1H), 4.38 – 4.26 (m, 1H), 2.28 (d, J = 9.9 Hz, 2H), 1.84 (d, J = 12.6 Hz, 2H), 1.54 – 1.40 (m, 1H), 1.40 – 1.30 (m, 2H), 1.30 – 1.14 (m, 2H), 0.98 (d, J = 6.5 Hz, 3H). ¹³C NMR (125 MHz, CDCl₃) δ 159.0, 158.8, 150.8, 150.5, 150.4, 135.7, 134.6, 132.7, 129.0, 125.7, 123.2, 120.5, 113.9, 50.4, 34.1, 33.1, 32.3, 22.3. HRMS (ESI): calcd for C₂₀H₂₃N₄ [M+H]⁺, 319.1917; found, 319.1921.

N-(4-Methoxycyclohexyl)-2-(pyridin-3-yl)quinazolin-4-amine (6d)—White solid (50 mg, 50%); mp 163–165 °C. ¹H NMR (500 MHz, CDCl₃) δ 9.73 (s, 1H), 8.81 (d, *J*=7.7

Hz, 1H), 8.70 (d, J = 3.4 Hz, 1H), 7.95 (d, J = 8.2 Hz, 1H), 7.81 – 7.57 (m, 2H), 7.52 – 7.32 (m, 2H), 5.69 (d, J = 0.8 Hz, 1H), 4.52 – 4.27 (m, 1H), 3.40 (s, 3H), 3.31 – 3.13 (m, 1H), 2.35 (d, J = 12.5 Hz, 2H), 2.18 (d, J = 11.3 Hz, 2H), 1.61 – 1.22 (m, 4H). ¹³C NMR (125 MHz, CDCl₃) δ 159.1, 158.7, 150.8, 150.3, 135.8, 134.3, 132.8, 128.9, 125.9, 123.3, 120.6, 113.8, 78.4, 56.0, 49.7, 30.6, 30.4. HRMS (ESI): calcd for C₂₀H₂₃N₄O [M+H]⁺, 335.1866; found, 335.1866.

N-(4,4-Dimethylcyclohexyl)-2-(pyridin-3-yl)quinazolin-4-amine (6e)—White solid (65 mg, 65%); mp 171–172 °C. ¹H NMR (500 MHz, CDCl₃) δ 9.74 (s, 1H), 8.80 (d, *J* = 7.9 Hz, 1H), 8.69 (d, *J* = 3.7 Hz, 1H), 7.90 (s, 1H), 7.79 – 7.63 (m, 2H), 7.45 (s, 1H), 7.41 (d, *J* = 3.1 Hz, 1H), 5.66 (d, *J* = 5.5 Hz, 1H), 4.43 – 4.25 (m, 1H), 2.16 – 2.00 (m, 2H), 1.64 – 1.38 (m, 6H), 1.01 (d, *J* = 3.0 Hz, 6H). ¹³C NMR (125 MHz, CDCl₃) δ 159.0, 158.8, 150.8, 150.4, 135.7, 134.5, 132.7, 129.0, 125.8, 123.2, 120.6, 113.9, 50.3, 37.9, 31.8, 29.9, 28.7, 25.2. HRMS (ESI): calcd for C₂₁H₂₅N₄ [M+H]⁺, 333.2074; found, 333.2076.

N-(4,4-Difluorocyclohexyl)-2-(pyridin-3-yl)quinazolin-4-amine (6f)—White solid (52 mg, 51%); mp 196–197 °C. ¹H NMR (500 MHz, CDCl₃) δ 9.72 (s, 1H), 8.80 (d, J= 7.8 Hz, 1H), 8.70 (d, J= 2.6 Hz, 1H), 7.95 (d, J= 8.3 Hz, 1H), 7.80 – 7.75 (m, 1H), 7.73 (d, J= 8.1 Hz, 1H), 7.47 (t, J= 7.5 Hz, 1H), 7.43 (dd, J= 7.7, 4.8 Hz, 1H), 5.70 (d, J= 6.2 Hz, 1H), 4.60 – 4.45 (m, 1H), 2.32 (d, J= 12.6 Hz, 2H), 2.28 – 2.12 (m, 2H), 2.12 – 1.93 (m, 2H), 1.85 – 1.70 (m, 2H). ¹³C NMR (125 MHz, CDCl₃) δ 159.1, 158.5, 150.8, 150.3, 150.1, 135.8, 134.3, 133.1, 129.0, 126.2, 123.6 (q, J_{C-F} = 241 Hz), 123.4, 120.5, 113.8, 48.1, 32.5 (t, J_{C-F} = 24.7 Hz), 28.6 (d, J_{C-F} = 9.2 Hz). HRMS (ESI): calcd for C₁₉H₁₉F₂N₄ [M+H]⁺, 341.1572; found, 341.1577.

N-(4-Dimethylaminocyclohexyl)-2-(pyridin-3-yl)quinazolin-4-amine (6g)—Paleyellow solid (63 mg, 60%); mp 231–235°C. ¹H NMR (500 MHz, CDCl₃) δ 9.70 (s, 1H), 8.76 (d, J = 7.8 Hz, 1H), 8.65 (d, J = 3.6 Hz, 1H), 7.87 (d, J = 8.2 Hz, 1H), 7.78 (d, J = 8.1 Hz, 1H), 7.75 – 7.66 (m, 1H), 7.42 (t, J = 7.5 Hz, 1H), 7.37 (dd, J = 7.5, 4.9 Hz, 1H), 6.02 (d, J = 6.7 Hz, 1H), 4.64 (s, 1H), 2.38 (s, 6H), 2.30 (s, 1H), 2.05 (s, 2H), 1.83 (s, 6H). ¹³C NMR (125 MHz, CDCl₃) δ 159.0, 158.6, 150.6, 150.3, 150.2, 135.7, 134.6, 132.7, 128.8, 125.9, 123.2, 120.8, 114.0, 62.2, 46.0, 42.5, 27.9, 25.6. HRMS (ESI): calcd for C₂₁H₂₆N₅ [M+H]⁺, 348.2183; found, 348.2184.

N-(Tetrahydro-2*H*-pyran-3-yl)-2-(pyridin-3-yl) quinazolin-4-amine (6h)—White solid (48 mg, 53%); mp 112–113 °C. ¹H NMR (500 MHz, CDCl₃) & 9.72 (s, 1H), 8.80 (dt, *J* = 7.9, 1.9 Hz, 1H), 8.68 (d, *J* = 3.7 Hz, 1H), 7.93 (d, *J* = 8.3 Hz, 1H), 7.80 (d, *J* = 8.2 Hz, 1H), 7.78 – 7.73 (m, 1H), 7.50 – 7.44 (m, 1H), 7.41 (dd, *J* = 7.8, 4.8 Hz, 1H), 6.18 (d, *J* = 7.1 Hz, 1H), 4.68 – 4.59 (m, 1H), 4.01 (dd, *J* = 11.5, 2.7 Hz, 1H), 3.89 – 3.80 (m, 1H), 3.80 – 3.66 (m, 2H), 2.08 – 2.00 (m, 3H), 1.95 – 1.83 (m, 1H), 1.73 – 1.61 (m, 1H). ¹³C NMR (125 MHz, CDCl₃) & 159.0, 158.6, 150.7, 150.3, 150.2, 135.9, 134.4, 133.0, 128.9, 126.1, 123.3, 120.7, 113.9, 71.5, 68.6, 46.5, 27.8, 23.0. HRMS (ESI): calcd for C₁₈H₁₉N₄O [M+H]⁺, 307.1553; found, 307.1556.

N-(1-Methylpiperidin-4-yl)-2-(pyridin-3-yl)quinazolin-4-amine (6i)—Pale-yellow solid (34 mg, 35%); mp 132–133 °C. ¹H NMR (500 MHz, CDCl₃) δ 9.72 (s, 1H), 8.78 (d, *J*

= 7.5 Hz, 1H), 8.68 (d, J = 3.4 Hz, 1H), 7.90 (d, J = 8.1 Hz, 1H), 7.73 (t, J = 7.4 Hz, 1H), 7.49 – 7.34 (m, 3H), 5.73 (d, J = 6.1 Hz, 1H), 4.48 – 4.30 (m, 1H), 2.91 (d, J = 9.1 Hz, 2H), 2.36 (s, 3H), 2.32 – 2.17 (m, 4H), 1.72 (m, 2H). ¹³C NMR (125 MHz, CDCl₃) & 159.0, 158.7, 150.8, 150.4, 150.3, 135.6, 134.5, 132.8, 129.0, 125.9, 123.2, 120.7, 113.9, 54.7, 48.0, 46.3, 32.1. HRMS (ESI): calcd for C₁₉H₂₂N₅ [M+H]⁺, 320.1870; found, 320.1872.

N-(1-Isopropylpiperidin-4-yl)-2-(pyridin-3-yl)quinazolin-4-amine (6j)—Yellow solid (49 mg, 47%); mp 191–192 °C. ¹H NMR (500 MHz, CDCl₃) δ 9.73 (d, *J* = 1.2 Hz, 1H), 8.79 (dt, *J* = 7.8, 1.5 Hz, 1H), 8.68 (dd, *J* = 4.6, 1.2 Hz, 1H), 7.91 (d, *J* = 8.3 Hz, 1H), 7.77 – 7.73 (m, 1H), 7.71 (d, *J* = 8.2 Hz, 1H), 7.49 – 7.43 (m, 1H), 7.41 (dd, *J* = 7.8, 4.8 Hz, 1H), 5.68 (d, *J* = 7.1 Hz, 1H), 4.49 – 4.35 (m, 1H), 2.99 (d, *J* = 11.1 Hz, 2H), 2.90 – 2.79 (m, 1H), 2.47 (t, *J* = 11.6 Hz, 2H), 2.28 (d, *J* = 11.5 Hz, 2H), 1.80 – 1.66 (m, 2H), 1.13 (d, *J* = 6.4 Hz, 6H). ¹³C NMR (125 MHz, CDCl₃) δ 159.0, 158.7, 150.8, 150.5, 150.4, 135.7, 134.5, 132.8, 129.0, 125.9, 123.3, 120.6, 113.9, 54.9, 48.7, 47.8, 32.3, 18.4. HRMS (ESI): calcd for C₂₁H₂₆N₅ [M+H]⁺, 348.2183; found, 348.2184.

N-(1-Benzylpiperidin-4-yl)-2-(pyridin-3-yl)quinazolin-4-amine (6k)—Yellow solid (67 mg, 57%); mp 180–182 °C. ¹H NMR (500 MHz, CDCl₃) & 9.72 (d, *J* = 1.6 Hz, 1H), 8.78 (dt, *J* = 7.9, 1.9 Hz, 1H), 8.69 (dd, *J* = 4.8, 1.6 Hz, 1H), 7.91 (d, *J* = 8.2 Hz, 1H), 7.77 – 7.72 (m, 1H), 7.71 (d, *J* = 8.1 Hz, 1H), 7.50 – 7.43 (m, 1H), 7.41 (dd, *J* = 7.8, 5.0 Hz, 1H), 7.39 – 7.27 (m, 5H), 5.66 (d, *J* = 7.1 Hz, 1H), 4.49 – 4.37 (m, 1H), 3.62 (s, 2H), 2.98 (d, *J* = 10.6 Hz, 2H), 2.38 – 2.29 (m, 2H), 2.24 (d, *J* = 11.4 Hz, 2H), 1.81 – 1.69 (m, 2H). ¹³C NMR (125 MHz, CDCl₃) & 159.0, 158.7, 150.8, 150.5, 150.3, 135.6, 134.5, 132.8, 129.4, 129.0, 128.4, 127.4, 125.9, 123.2, 120.6, 113.9, 63.2, 52.5, 48.4, 32.0. HRMS (ESI): calcd for C₂₅H₂₆N₅ [M+H]⁺, 396.2183; found, 396.2184.

N-Cyclooctyl-2-(pyridin-3-yl)quinazolin-4-amine (7a)—White solid (65 mg, 66%); mp 140–142 °C. ¹H NMR (500 MHz, CDCl₃) & 9.75 (d, J= 1.4 Hz, 1H), 8.82 (d, J= 7.9 Hz, 1H), 8.69 (dd, J= 4.7, 1.4 Hz, 1H), 7.92 (d, J= 8.3 Hz, 1H), 7.76 – 7.72 (m, 1H), 7.70 (d, J= 8.1 Hz, 1H), 7.51 – 7.35 (m, 2H), 5.73 (d, J= 6.2 Hz, 1H), 4.73 – 4.56 (m, 1H), 2.18 – 2.00 (m, 2H), 2.00 – 1.43 (m, 12H). ¹³C NMR (125 MHz, CDCl₃) & 158.8, 158.6, 150.7, 150.3, 150.2, 135.8, 134.5, 132.7, 128.9, 125.8, 123.3, 120.5, 114.0, 51.0, 32.7, 27.2, 26.1, 24.2. HRMS (ESI): calcd for C₂₁H₂₅N₄ [M+H]⁺, 333.2074; found, 333.2076.

N-Cycloheptyl-2-(pyridin-3-yl)quinazolin-4-amine (7b)—Off-white solid (70 mg, 74%); mp 158–159°C. ¹H NMR (500 MHz, CDCl₃) δ 9.75 (s, 1H), 8.82 (d, *J* = 7.9 Hz, 1H), 8.68 (s, 1H), 7.93 (d, *J* = 8.3 Hz, 1H), 7.78 – 7.65 (m, 2H), 7.50 – 7.37 (m, 2H), 5.76 (d, *J* = 5.6 Hz, 1H), 4.64 – 4.46 (m, 1H), 2.25 – 2.12 (m, 2H), 1.83 – 1.52 (m, 10H). ¹³C NMR (125 MHz, CDCl₃) δ 158.7, 158.6, 150.7, 150.3, 150.2, 135.8, 134.5, 132.7, 128.8, 125.9, 123.3, 120.6, 113.9, 52.3, 34.9, 28.3, 24.6. HRMS (ESI): calcd for C₂₀H₂₃N₄ [M+H]⁺, 319.1917; found, 319.1918.

N-Cyclopentyl-2-(pyridin-3-yl)quinazolin-4-amine (7c)—Off-white solid (52 mg, 60%); mp 147–148°C. ¹H NMR (500 MHz, CDCl₃) δ 9.76 (d, *J* = 1.5 Hz, 1H), 8.81 (dt, *J* = 7.9, 1.9 Hz, 1H), 8.68 (dd, *J* = 4.8, 1.7 Hz, 1H), 7.91 (d, *J* = 8.2 Hz, 1H), 7.76 – 7.71 (m, 1H), 7.70 (d, *J* = 8.2 Hz, 1H), 7.46 – 7.42 (m, 1H), 7.40 (dd, *J* = 7.9, 4.8 Hz, 1H), 5.73 (d, *J*

= 6.1 Hz, 1H), 4.80 – 4.72 (m, 1H), 2.34 – 2.25 (m, 2H), 1.86 – 1.71 (m, 4H), 1.67 – 1.59 (m, 2H). 13 C NMR (125 MHz, CDCl₃) δ 159.4, 158.8, 150.7, 150.4, 150.3, 135.7, 134.6, 132.7, 129.0, 125.8, 123.2, 120.6, 113.9, 53.2, 33.4, 24.1. HRMS (ESI): calcd for C₁₈H₁₉N₄ [M+H]⁺, 291.1604; found, 291.1604.

N-Cyclobutyl-2-(pyridin-3-yl)quinazolin-4-amine (7d)—Off-white solid (47 mg, 57%); mp 173–174°C. ¹H NMR (500 MHz, CDCl₃) & 9.75 (s, 1H), 8.87 – 8.79 (m, 1H), 8.69 (d, *J* = 4.2 Hz, 1H), 7.93 (dd, *J* = 8.0, 4.1 Hz, 1H), 7.74 (t, *J* = 8.2 Hz, 2H), 7.47 – 7.43 (m, 1H), 7.43 – 7.39 (m, 1H), 6.03 (s, 1H), 4.96 – 4.97 (m, 1H), 2.66 – 2.54 (m, 2H), 2.16 – 2.03 (m, 2H), 1.96 – 1.85 (m, 2H). ¹³C NMR (125 MHz, CDCl₃) & 158.8, 150.7, 150.5, 150.4, 135.7, 134.5, 132.8, 129.0, 125.9, 123.2, 120.6, 113.7, 46.8, 31.4, 15.6. HRMS (ESI): calcd for C₁₇H₁₇N₄ [M+H]⁺, 277.1448; found, 277.1449.

N-Cyclopropyl-2-(pyridin-3-yl)quinazolin-4-amine (7e)—Pale-yellow solid (30 mg, 38%); mp 180–181°C. ¹H NMR (500 MHz, CDCl₃) δ 9.81 (s, 1H), 8.89 (d, *J* = 7.9 Hz, 1H), 8.70 (d, *J* = 3.7 Hz, 1H), 7.96 (d, *J* = 8.3 Hz, 1H), 7.77 – 7.73 (m, 1H), 7.71 (d, *J* = 8.2 Hz, 1H), 7.47 – 7.36 (m, 2H), 6.14 (s, 1H), 3.17 (td, *J* = 6.9, 3.1 Hz, 1H), 1.07 – 0.95 (m, 2H), 0.79 – 0.67 (m, 2H). ¹³C NMR (125 MHz, CDCl₃) δ 160.9, 158.9, 150.8, 150.5, 150.4, 135.8, 134.5, 132.8, 129.0, 126.0, 123.2, 120.6, 113.9, 24.5, 7.5. HRMS (ESI): calcd for C₁₆H₁₅N₄ [M+H]⁺, 263.1291; found, 263.1291.

N-(Adamantan-1-yl)-2-(pyridin-3-yl)quinazolin-4-amine (7f)—White solid (37 mg, 35%); mp 114–116°C. ¹H NMR (500 MHz, CDCl₃) δ 9.73 (d, *J* = 1.6 Hz, 1H), 8.83 (d, *J* = 7.8 Hz, 1H), 8.69 (dd, *J* = 4.7, 1.5 Hz, 1H), 7.92 (d, *J* = 8.1 Hz, 1H), 7.80 – 7.69 (m, 1H), 7.66 (d, *J* = 8.1 Hz, 1H), 7.51 – 7.32 (m, 2H), 5.58 (s, 1H), 2.38 (d, *J* = 2.3 Hz, 6H), 2.25 – 2.18 (m, 3H), 1.88 – 1.74 (m, 6H). ¹³C NMR (125 MHz, CDCl₃) δ 159.0, 158.1, 150.7, 150.3, 135.8, 134.5, 132.6, 128.9, 125.9, 123.3, 120.6, 114.2, 53.6, 41.7, 36.7, 29.7. HRMS (ESI): calcd for C₂₃H₂₅N₄ [M+H]⁺, 357.2074; found, 357.2074.

N-(Bicyclo[2.2.1]heptan-2-yl)-2-(pyridin-3-yl)quinazolin-4-amine (7g)—White solid (47 mg, 50%); mp 145–146°C. ¹H NMR (500 MHz, CDCl₃) & 9.75 (s, 1H), 8.82 (dt, J = 7.9, 1.8 Hz, 1H), 8.68 (d, J = 3.9 Hz, 1H), 7.91 (d, J = 8.3 Hz, 1H), 7.75 – 7.71 (m, 1H), 7.69 (d, J = 8.2 Hz, 1H), 7.49 – 7.35 (m, 2H), 5.64 (d, J = 5.5 Hz, 1H), 4.27 – 4.16 (m, 1H), 2.51 (d, J = 4.1 Hz, 1H), 2.39 (s, 1H), 2.05 (ddd, J = 13.2, 7.9, 2.2 Hz, 1H), 1.73 – 1.61 (m, 1H), 1.61 – 1.53 (m, 1H), 1.51 (d, J = 10.1 Hz, 1H), 1.48 – 1.36 (m, 2H), 1.34 – 1.20 (m, 2H). ¹³C NMR (125 MHz, CDCl₃) & 158.9, 158.7, 150.7, 150.3, 150.2, 135.8, 134.5, 132.7, 128.9, 125.8, 123.3, 120.5, 113.9, 55.1, 42.1, 41.0, 36.0, 35.9, 28.3, 26.7. HRMS (ESI): calcd for C₂₀H₂₁N₄ [M+H]⁺, 317.1761; found, 317.1763.

N-(Cyclohexylmethyl)-2-(pyridin-3-yl)quinazolin-4-amine (7h)—Pale-yellow solid (67 mg, 71%); mp 121–123°C. ¹H NMR (500 MHz, CDCl₃) & 9.75 (s, 1H), 8.81 (d, *J* = 7.8 Hz, 1H), 8.68 (d, *J* = 3.8 Hz, 1H), 7.91 (d, *J* = 8.2 Hz, 1H), 7.79 – 7.65 (m, 2H), 7.50 – 7.34 (m, 2H), 5.98 (s, 1H), 3.65 (t, *J* = 6.1 Hz, 2H), 1.87 (d, *J* = 12.6 Hz, 2H), 1.82 – 1.61 (m, 4H), 1.32 – 1.18 (m, 3H), 1.14 – 1.04 (m, 1H). ¹³C NMR (125 MHz, CDCl₃) & 159.9, 158.8, 150.7, 150.3, 135.7, 134.6, 132.7, 128.9, 125.8, 123.2, 120.5, 114.0, 47.6, 38.0, 31.3, 26.5, 26.0. HRMS (ESI): calcd for C₂₀H₂₃N₄ [M+H]⁺, 319.1917; found, 319.1920.

N-(2-Cyclohexylethyl)-2-(pyridin-3-yl)quinazolin-4-amine (7i)—White solid (48 mg, 48%); mp 153–154°C. ¹H NMR (500 MHz, CDCl₃) δ 9.76 (s, 1H), 8.86 – 8.78 (m, 1H), 8.69 (d, *J* = 4.1 Hz, 1H), 7.92 (dd, *J* = 7.9, 5.4 Hz, 1H), 7.79 – 7.67 (m, 2H), 7.48 – 7.37 (m, 2H), 5.84 (s, 1H), 3.86 – 3.75 (m, 2H), 1.83 (d, *J* = 12.3 Hz, 2H), 1.76 – 1.60 (m, 5H), 1.50 – 1.38 (m, 1H), 1.33 – 1.12 (m, 3H), 1.08 – 0.96 (m, 2H). ¹³C NMR (125 MHz, CDCl₃) δ 159.7, 158.8, 150.7, 150.4, 150.3, 135.8, 135.7, 134.5, 132.8, 132.7, 128.9, 128.9, 125.9, 123.2, 120.6, 114.0, 39.4, 37.0, 35.7, 33.4, 26.6, 26.3. HRMS (ESI): calcd for C₂₁H₂₅N₄ [M+H]⁺, 333.2074; found, 333.2074.

N-Phenyl-2-(pyridin-3-yl)quinazolin-4-amine (8a)—Off-white solid (55 mg, 62%); mp 179–180°C. ¹H NMR (500 MHz, CDCl₃) & 9.73 (d, J= 1.8 Hz, 1H), 8.82 (dt, J= 7.9, 1.9 Hz, 1H), 8.70 (dd, J= 4.8, 1.6 Hz, 1H), 8.01 (d, J= 8.3 Hz, 1H), 7.95 (d, J= 8.1 Hz, 1H), 7.86 (d, J= 7.6 Hz, 2H), 7.84 – 7.80 (m, 1H), 7.69 (brs, 1H), 7.61 – 7.52 (m, 1H), 7.50 – 7.45 (m, 2H), 7.43 (dd, J= 7.9, 4.8 Hz, 1H), 7.21 (t, J= 7.4 Hz, 1H). ¹³C NMR (125 MHz, CDCl₃) & 158.4, 157.6, 150.8, 150.7, 150.2, 138.4, 136.1, 134.2, 133.3, 129.3, 129.2, 126.8, 124.6, 123.4, 121.6, 120.5, 114.1. HRMS (ESI): calcd for C₁₉H₁₅N₄ [M+H]⁺, 299.1291; found, 299.1292.

N-Benzyl-2-(pyridin-3-yl)quinazolin-4-amine (8b)—Off-white solid (65 mg, 70%); mp 140–141°C. ¹H NMR (500 MHz, CDCl₃) δ 9.56 (s, 1H), 8.61 (d, *J* = 7.9 Hz, 1H), 8.47 (d, *J* = 3.6 Hz, 1H), 7.74 (d, *J* = 8.3 Hz, 1H), 7.61 (d, *J* = 8.2 Hz, 1H), 7.58 – 7.44 (m, 1H), 7.28 – 7.18 (m, 4H), 7.18 – 7.13 (m, 2H), 7.13 – 7.07 (m, 1H), 6.39 (s, 1H), 4.78 (d, *J* = 5.4 Hz, 2H). ¹³C NMR (125 MHz, CDCl₃) δ 159.6, 158.7, 150.6, 150.4, 150.2, 138.5, 135.8, 134.5, 132.8, 128.0, 127.7, 126.0, 123.2, 120.9, 113.9, 45.4. HRMS (ESI): calcd for C₂₀H₁₇N₄ [M+H]⁺, 313.1448; found, 313.1453.

N-Phenethyl-2-(pyridin-3-yl)quinazolin-4-amine (8c)—White solid (43 mg, 44%); mp 140–141°C. ¹H NMR (500 MHz, CDCl₃) δ 9.79 (s, 1H), 8.85 (d, *J* = 7.1 Hz, 1H), 8.70 (s, 1H), 7.94 (d, *J* = 8.4 Hz, 1H), 7.78 – 7.69 (m, 1H), 7.60 (d, *J* = 8.0 Hz, 1H), 7.46 – 7.39 (m, 2H), 7.38 – 7.31 (m, 2H), 7.30 – 7.25 (m, 3H), 5.98 (s, 1H), 4.11 – 3.99 (m, 2H), 3.11 (t, *J* = 6.9 Hz, 2H). ¹³C NMR (125 MHz, CDCl₃) δ 159.6, 158.7, 150.8, 150.3, 150.2, 139.0, 135.8, 134.4, 132.9, 129.0, 128.9, 126.8, 126.0, 123.3, 120.5, 113.9, 42.6, 35.4. HRMS (ESI): calcd for C₂₁H₁₉N₄ [M+H]⁺, 327.1604; found, 327.1603.

N-(3-Phenylpropyl)-2-(pyridin-3-yl)quinazolin-4-amine (8d)—Off-white sticky oil (60 mg, 59%). ¹H NMR (500 MHz, CDCl₃) δ 9.74 (s, 1H), 8.76 (dt, *J* = 7.9, 1.8 Hz, 1H), 8.68 (d, *J* = 3.6 Hz, 1H), 7.90 (d, *J* = 8.3 Hz, 1H), 7.78 – 7.66 (m, 1H), 7.53 (d, *J* = 7.8 Hz, 1H), 7.45 – 7.34 (m, 2H), 7.33 – 7.27 (m, 2H), 7.25 – 7.19 (m, 3H), 5.94 (s, 1H), 3.89 – 3.75 (m, 2H), 2.81 (t, *J* = 7.4 Hz, 2H), 2.13 (p, *J* = 7.2 Hz, 2H). ¹³C NMR (125 MHz, CDCl₃) δ 159.7, 158.7, 150.7, 150.3, 150.2, 141.6, 135.8, 134.5, 132.7, 128.8, 128.7, 128.5, 126.2, 125.8, 123.2, 120.6, 113.9, 41.2, 33.7, 30.7. HRMS (ESI): calcd for C₂₂H₂₁N₄ [M+H]⁺, 341.1761; found, 341.1758.

N-(4-Phenylbutyl)-2-(pyridin-3-yl)quinazolin-4-amine (8e)—Pale-yellow solid (73 mg, 69%); mp 108–109°C. ¹H NMR (500 MHz, CDCl₃) δ 9.75 (s, 1H), 8.81 (d, *J* = 7.9 Hz, 1H), 8.69 (d, *J* = 3.7 Hz, 1H), 7.93 (d, *J* = 8.3 Hz, 1H), 7.77 – 7.71 (m, 1H), 7.70 (d, *J* = 8.1

Hz, 1H), 7.51 - 7.35 (m, 2H), 7.28 (d, J = 7.7 Hz, 2H), 7.23 - 7.10 (m, 3H), 5.90 (s, 1H), 3.89 - 3.70 (m, 2H), 2.72 (t, J = 6.7 Hz, 2H), 1.90 - 1.75 (m, 4H). ¹³C NMR (125 MHz, CDCl₃) δ 159.7, 158.7, 150.8, 150.3, 150.1, 142.1, 135.8, 134.4, 132.8, 128.8, 128.5, 128.6, 125.9, 123.3, 120.6, 113.9, 41.4, 35.7, 29.0, 28.9. HRMS (ESI): calcd for C₂₃H₂₃N₄ [M+H]⁺, 355.1917; found, 355.1919.

N-(2-(Pyridin-2-yl)ethyl)-2-(pyridin-3-yl)quinazolin-4-amine (8f)—White solid (57 mg, 58%); mp 138–139°C. ¹H NMR (500 MHz, CDCl₃) & 9.75 (d, *J* = 1.2 Hz, 1H), 8.82 (dt, *J* = 7.9, 1.9 Hz, 1H), 8.70 – 8.66 (m, 1H), 8.51 (s, 1H), 8.47 (d, *J* = 4.1 Hz, 1H), 7.93 (d, *J* = 8.2 Hz, 1H), 7.77 – 7.72 (m, 1H), 7.68 (d, *J* = 8.1 Hz, 1H), 7.59 (d, *J* = 7.8 Hz, 1H), 7.46 – 7.38 (m, 2H), 7.24 (dd, *J* = 7.7, 4.9 Hz, 1H), 6.28 (t, *J* = 5.3 Hz, 1H), 4.10 – 3.96 (m, 2H), 3.13 (t, *J* = 7.0 Hz, 2H). ¹³C NMR (125 MHz, CDCl₃) & 159.7, 158.6, 150.7, 150.2, 150.1, 148.1, 136.6, 135.8, 134.7, 134.4, 133.0, 128.9, 126.2, 123.8, 123.4, 120.7, 113.9, 42.4, 32.6. HRMS (ESI): calcd for C₂₀H₁₈N₅ [M+H]⁺, 328.1557; found, 328.1561.

N-(2-(Pyridin-3-yl)ethyl)-2-(pyridin-3-yl)quinazolin-4-amine (8g)—Off-white solid (67 mg, 68%); mp 102–104°C. ¹H NMR (500 MHz, CDCl₃) δ 9.77 (d, J= 1.4 Hz, 1H), 8.83 (dt, J= 7.9, 1.9 Hz, 1H), 8.68 (dd, J= 4.8, 1.7 Hz, 1H), 7.91 (d, J= 8.0 Hz, 1H), 7.73 (ddd, J = 8.3, 7.0, 1.3 Hz, 1H), 7.57 (d, J= 7.8 Hz, 1H), 7.46 – 7.37 (m, 2H), 7.07 (d, J= 8.3 Hz, 2H), 6.72 – 6.62 (m, 2H), 5.96 – 5.87 (m, 1H), 4.03 – 3.94 (m, 2H), 2.98 (t, J= 6.9 Hz, 2H). ¹³C NMR (125 MHz, CDCl₃) δ 159.6, 158.8, 150.7, 150.3, 145.1, 135.8, 134.5, 132.8, 129.8, 128.9, 128.8, 125.9, 123.3, 120.6, 115.6, 114.0, 42.8, 34.5. HRMS (ESI): calcd for C₂₀H₁₈N₅ [M+H]⁺, 328.1557; found, 328.1560.

(S)-2-(Pyridin-3-yl)-N-(1,2,3,4-tetrahydronaphthalen-2-yl)quinazolin-4-amine

(9a)—Off-white solid (74 mg, 70%); mp 150–152°C. ¹H NMR (500 MHz, CDCl₃) & 9.75 (s, 1H), 8.82 (d, *J* = 7.9 Hz, 1H), 8.68 (d, *J* = 3.7 Hz, 1H), 7.95 (d, *J* = 8.3 Hz, 1H), 7.75 (t, *J* = 7.7 Hz, 1H), 7.70 (d, *J* = 8.1 Hz, 1H), 7.47 – 7.42 (m, 1H), 7.41 (dd, *J* = 7.9, 4.8 Hz, 1H), 7.24 – 7.11 (m, 4H), 5.87 (d, *J* = 5.2 Hz, 1H), 5.00 – 4.87 (m, 1H), 3.43 (dd, *J* = 16.2, 5.0 Hz, 1H), 3.12 – 2.88 (m, 3H), 2.40 – 2.30 (m, 1H), 2.13 – 2.01 (m, 1H). ¹³C NMR (125 MHz, CDCl₃) & 159.2, 158.7, 150.8, 150.2, 135.8, 135.7, 134.3, 134.1, 132.9, 129.7, 129.1, 128.9, 126.4, 126.2, 126.0, 123.3, 120.7, 113.9, 47.1, 35.8, 28.5, 27.5. HRMS (ESI): calcd for C₂₃H₂₁N₄ [M+H]⁺, 353.1761; found, 328.1762.

(R)-2-(Pyridin-3-yl)-N-(1,2,3,4-tetrahydronaphthalen-2-yl)quinazolin-4-amine

(9b)—Off-white solid (65 mg, 62%); mp 153–156°C. ¹H NMR (500 MHz, CDCl₃) & 9.75 (s, 1H), 8.82 (d, *J* = 7.9 Hz, 1H), 8.68 (d, *J* = 3.7 Hz, 1H), 7.95 (d, *J* = 8.3 Hz, 1H), 7.75 (t, *J* = 7.7 Hz, 1H), 7.70 (d, *J* = 8.1 Hz, 1H), 7.47 – 7.42 (m, 1H), 7.41 (dd, *J* = 7.9, 4.8 Hz, 1H), 7.24 – 7.11 (m, 4H), 5.87 (d, *J* = 5.2 Hz, 1H), 5.00 – 4.87 (m, 1H), 3.43 (dd, *J* = 16.2, 5.0 Hz, 1H), 3.12 – 2.88 (m, 3H), 2.40 – 2.30 (m, 1H), 2.13 – 2.01 (m, 1H). ¹³C NMR (125 MHz, CDCl₃) & 159.2, 158.7, 150.8, 150.2, 135.8, 135.7, 134.3, 134.1, 132.9, 129.7, 129.1, 128.9, 126.4, 126.2, 126.0, 123.3, 120.7, 113.9, 47.1, 35.8, 28.5, 27.5. HRMS (ESI): calcd for C₂₃H₂₁N₄ [M+H]⁺, 353.1761; found, 328.1764.

N-(2,3-Dihydro-1*H*-inden-2-yl)-2-(pyridin-3-yl)quinazolin-4-amine (9c)—Grey solid (75 mg, 74%); mp 167–168°C. ¹H NMR (500 MHz, CDCl₃) δ 9.78 (d, *J* = 1.7 Hz,

1H), 8.84 (dd, J= 7.9, 1.7 Hz, 1H), 8.69 (d, J= 4.7 Hz, 1H), 7.93 (d, J= 8.3 Hz, 1H), 7.77 – 7.70 (m, 1H), 7.68 (d, J= 8.1 Hz, 1H), 7.47 – 7.37 (m, 2H), 7.32 – 7.27 (m, 2H), 7.25 – 7.20 (m, 2H), 6.04 (s, 1H), 5.37 – 5.29 (m, 1H), 3.60 (dd, J= 16.2, 7.2 Hz, 2H), 3.08 (dd, J= 16.2, 4.8 Hz, 2H). ¹³C NMR (125 MHz, CDCl₃) δ 159.4, 158.6, 150.7, 150.3, 141.1, 135.8, 134.4, 132.9, 128.9, 127.0, 126.0, 125.1, 123.3, 120.7, 113.9, 52.6, 40.3. HRMS (ESI): calcd for C₂₄H₂₁NO [M+H]⁺, 339.1618; found, 339.1612.

N-(1,2,3,4-Tetrahydronaphthalen-1-yl)-2-(pyridin-3-yl) quinazolin-4-amine (9d)

—Pale-yellow sticky oil (35 mg, 33%). ¹H NMR (500 MHz, CDCl₃) δ 9.78 (d, J= 0.9 Hz, 1H), 8.89 (d, J= 7.8 Hz, 1H), 8.69 (d, J= 3.5 Hz, 1H), 8.00 (d, J= 7.8 Hz, 1H), 7.83 – 7.73 (m, 1H), 7.68 (d, J= 8.0 Hz, 1H), 7.51 – 7.41 (m, 2H), 7.39 (d, J= 7.6 Hz, 1H), 7.25 – 7.22 (m, 1H), 7.22 – 7.15 (m, 2H), 6.02 (s, 1H), 5.94 – 5.86 (m, 1H), 3.00 – 2.80 (m, 2H), 2.34 – 2.23 (m, 1H), 2.20 – 2.12 (m, 1H), 2.00 – 1.93 (m, 2H). ¹³C NMR (125 MHz, CDCl₃) δ 159.0, 158.7, 150.6, 150.2, 138.1, 136.8, 136.1, 133.0, 129.5, 129.1, 128.8, 127.7, 126.6, 126.1, 123.4, 120.7, 113.8, 49.2, 29.6, 29.5, 20.2. HRMS (ESI): calcd for C₂₃H₂₁N₄ [M +H]⁺, 353.1761; found, 353.1767.

N-(2,3-Dihydro-1*H*-inden-1-yl)-2-(pyridin-3-yl)quinazolin-4-amine (9e)—Brown solid (57 mg, 56%); mp 128–130°C. ¹H NMR (500 MHz, CDCl₃) & 9.78 (d, J= 1.5 Hz, 1H), 8.84 (dt, J= 7.9, 1.9 Hz, 1H), 8.67 (dd, J= 4.8, 1.6 Hz, 1H), 7.95 (d, J= 8.4 Hz, 1H), 7.80 – 7.73 (m, 1H), 7.70 (d, J= 8.0 Hz, 1H), 7.50 – 7.37 (m, 3H), 7.35 (d, J= 7.5 Hz, 1H), 7.33 – 7.28 (m, 1H), 7.26 – 7.20 (m, 2H), 6.17 (q, J= 7.4 Hz, 1H), 6.02 (d, J= 7.5 Hz, 1H), 3.12 (ddd, J= 15.8, 8.7, 4.0 Hz, 1H), 3.04 (dt, J= 16.0, 8.0 Hz, 1H), 2.92 – 2.83 (m, 1H), 2.16 – 2.01 (m, 1H). ¹³C NMR (125 MHz, CDCl₃) & 159.5, 158.8, 150.8, 150.5, 150.3, 143.9, 143.4, 135.8, 134.5, 132.9, 129.0, 128.4, 127.0, 126.0, 125.2, 124.3, 123.3, 120.7, 113.8, 56.5, 34.1, 30.5. HRMS (ESI): calcd for C₂₂H₁₉N₄ [M+H]⁺, 339.1604; found, 339.1610.

N-(Chroman-3-yl)-2-(pyridin-3-yl)quinazolin-4-amine (9f)—White solid (67 mg, 63%); mp 145–148°C. ¹H NMR (500 MHz, CDCl₃) δ 9.74 (s, 1H), 8.80 (dt, J= 7.9, 1.9 Hz, 1H), 8.68 (d, J= 3.6 Hz, 1H), 7.90 (d, J= 7.9 Hz, 1H), 7.72 (ddd, J= 8.3, 7.0, 1.3 Hz, 1H), 7.63 (d, J= 7.7 Hz, 1H), 7.45 – 7.34 (m, 2H), 7.19 – 7.12 (m, 1H), 7.07 (d, J= 6.6 Hz, 1H), 6.95 – 6.88 (m, 2H), 6.09 (d, J= 7.3 Hz, 1H), 5.11 (ddt, J= 7.3, 3.6, 1.7 Hz, 1H), 4.43 (ddd, J= 11.0, 3.8, 2.2 Hz, 1H), 4.35 (dd, J= 11.0, 1.7 Hz, 1H), 3.34 (dd, J= 16.8, 5.2 Hz, 1H), 3.10 (d, J= 16.8 Hz, 1H). ¹³C NMR (125 MHz, CDCl₃) δ 159.0, 158.4, 153.9, 150.7, 150.3, 150.1, 135.6, 134.2, 132.9, 130.6, 128.9, 127.8, 126.0, 123.2, 121.5, 120.7, 119.5, 117.0, 113.7, 68.0, 44.0, 30.4. HRMS (ESI): calcd for C₂₂H₁₉N₄O [M+H]⁺, 355.1553; found, 355.1551.

Preparation of 2-chloro-*N***-(2,3-dihydro-1***H***-inden-2-yl)quinazolin-4-amine (10)**— A mixture of 2,4-dichloroquinazoline (398 mg, 2.0 mmol), 2,3-dihydro-1*H*-inden-2-amine (266 mg, 2.0 mmol), and potassium carbonate (276 mg, 2.0 mmol) in DMF (5 mL) was stirred at room temperature for 5 h. Water (20 mL) was added, and the formed solid was filtered, washed with water, and solid was dried to give 10 as an off-white solid (390 mg, 66 %); mp 239–240°C.

¹H NMR (500 MHz, CDCl₃) δ 7.76 (d, J= 8.2 Hz, 1H), 7.74 – 7.69 (m, 1H), 7.60 (d, J= 8.2 Hz, 1H), 7.41 (t, J= 7.4 Hz, 1H), 7.30 – 7.26 (m, 2H), 7.24 – 7.19 (m, 2H), 6.08 (d, J= 6.7 Hz, 1H), 5.27 – 5.15 (m, 1H), 3.53 (dd, J= 16.2, 7.0 Hz, 2H), 3.00 (dd, J= 16.2, 4.0 Hz, 2H). ¹³C NMR (125 MHz, CDCl₃) δ 160.6, 157.8, 151.0, 140.7, 133.6, 128.0, 127.1, 126.2, 125.1, 120.8, 113.3, 52.6, 40.2. MS (ESI) m/z [M + H]+: calcd, 296.09; found, 296.13.

General Procedure for 11a-11h—A mixture of 2-chloro-*N*-(2,3-dihydro-1*H*-inden-2yl)quinazolin-4-amine **10** (148 mg, 0.5 mmol), boronic acid (0.5 mmol), Pd(PPh₃)₄ (58 mg, 0.05 mmol), potassium carbonate (276 mg, 2.0 mmol) in dioxane (10 mL) and water (1.5 mL) was heated at 85 °C under an argon atmosphere for 16 h. Water (5 mL) was added, and the mixture was extracted with EtOAc (25 mL × 3). The combined organic phase was washed with brine (15 mL), dried (Na₂SO₄), filtered, evaporated, and purified by flash chromatography to give product.

Analytical Data for Compounds 11a-11h

N-(2,3-Dihydro-1*H*-inden-2-yl)-2-(4-methylpyridin-3-yl)quinazolin-4-amine (11a) —White solid (125 mg, 71%); mp 170–172°C. ¹H NMR (500 MHz, CDCl₃) δ 9.16 (s, 1H), 8.50 (d, *J* = 5.0 Hz, 1H), 7.91 (d, *J* = 7.9 Hz, 1H), 7.75 (ddd, *J* = 8.3, 7.0, 1.2 Hz, 1H), 7.67 (d, *J* = 7.8 Hz, 1H), 7.49 – 7.42 (m, 1H), 7.29 (dd, *J* = 5.3, 3.4 Hz, 2H), 7.25 – 7.19 (m, 3H), 5.94 (d, *J* = 7.0 Hz, 1H), 5.26 (tdd, *J* = 7.1, 4.4, 2.6 Hz, 1H), 3.54 (dd, *J* = 16.2, 7.1 Hz, 2H), 3.04 (dd, *J* = 16.2, 4.4 Hz, 2H), 2.72 (s, 3H). ¹³C NMR (125 MHz, CDCl₃) δ 161.1, 159.1, 151.3, 150.1, 149.2, 146.6, 140.9, 135.4, 132.7, 128.8, 126.8, 125.9, 124.9, 120.7, 113.2, 52.3, 40.2, 21.0. HRMS (ESI): calcd for C₂₃H₂₁N₄ [M+H]⁺, 353.1761; found, 353.1765.

N-(2,3-Dihydro-1*H*-inden-2-yl)-2-(5-methylpyridin-3-yl)quinazolin-4-amine (11b) —Yellow solid (134 mg, 76%); mp 200–201°C. ¹H NMR (500 MHz, CDCl₃) & 9.59 (d, J= 1.6 Hz, 1H), 8.63 (s, 1H), 8.53 (d, J= 1.6 Hz, 1H), 7.92 (d, J= 8.0 Hz, 1H), 7.77 – 7.71 (m, 1H), 7.66 (d, J= 7.8 Hz, 1H), 7.45 – 7.38 (m, 1H), 7.30 (dd, J= 5.3, 3.4 Hz, 2H), 7.23 (dd, J= 5.5, 3.2 Hz, 2H), 5.95 (d, J= 6.8 Hz, 1H), 5.40 – 5.29 (m, 1H), 3.61 (dd, J= 16.2, 7.2 Hz, 2H), 3.08 (dd, J= 16.2, 4.8 Hz, 2H), 2.45 (s, 3H). ¹³C NMR (125 MHz, CDCl₃) & 159.3, 158.8, 151.2, 150.3, 147.6, 141.1, 136.0, 134.0, 132.7, 132.6, 128.7, 126.8, 125.7, 124.9, 120.8, 113.9, 52.4, 40.1, 18.5. HRMS (ESI): calcd for C₂₃H₂₁N₄ [M+H]⁺, 353.1761; found, 353.1757.

N-(2,3-Dihydro-1*H***-inden-2-yl)-2-(6-methylpyridin-3-yl)quinazolin-4-amine (11c)** —Yellow solid (129 mg, 73%); mp 175–176°C. ¹H NMR (500 MHz, CDCl₃) δ 9.66 (d, *J* = 1.9 Hz, 1H), 8.69 (dd, *J* = 8.1, 2.2 Hz, 1H), 7.89 (d, *J* = 8.0 Hz, 1H), 7.73 – 7.68 (m, 1H), 7.63 (d, *J* = 7.9 Hz, 1H), 7.41 – 7.36 (m, 1H), 7.29 – 7.25 (m, 2H), 7.23 – 7.19 (m, 3H), 5.91

(d, J = 6.6 Hz, 1H), 5.33 - 5.23 (m, 1H), 3.58 (dd, J = 16.2, 7.2 Hz, 2H), 3.06 (dd, J = 16.1, 4.8 Hz, 2H), 2.62 (s, 3H). ¹³C NMR (125 MHz, CDCl₃) δ 159.8, 159.2, 158.8, 150.4, 149.7, 141.0, 136.0, 132.6, 131.6, 128.8, 126.9, 125.6, 124.9, 122.7, 120.5, 113.7, 52.5, 40.2, 24.5. HRMS (ESI): calcd for C₂₃H₂₁N₄ [M+H]⁺, 353.1761; found, 353.1767.

N-(2,3-Dihydro-1*H*-inden-2-yl)-2-phenylquinazolin-4-amine (11d)—White solid (84 mg, 50%); mp 221–223°C. ¹H NMR (500 MHz, CDCl₃) δ 8.64 – 8.56 (m, 2H), 7.93 (d,

 $J=8.3 \text{ Hz}, 1\text{H}, 7.72 \text{ (ddd, } J=8.3, 7.0, 1.3 \text{ Hz}, 1\text{H}), 7.64 \text{ (d, } J=7.8 \text{ Hz}, 1\text{H}), 7.53 - 7.46 \text{ (m, 3H)}, 7.39 \text{ (ddd, } J=8.1, 7.0, 1.1 \text{ Hz}, 1\text{H}), 7.31 \text{ (dd, } J=5.3, 3.3 \text{ Hz}, 2\text{H}), 7.24 \text{ (dd, } J=5.5, 3.2 \text{ Hz}, 2\text{H}), 5.86 \text{ (d, } J=6.6 \text{ Hz}, 1\text{H}), 5.38 \text{ (tdd, } J=7.1, 5.1, 2.1 \text{ Hz}, 1\text{H}), 3.61 \text{ (dd, } J=16.1, 7.2 \text{ Hz}, 2\text{H}), 3.08 \text{ (dd, } J=16.1, 5.0 \text{ Hz}, 2\text{H}). {}^{13}\text{C} \text{ NMR} (125 \text{ MHz}, \text{CDCl}_3) \\ 8 \text{ 160.4}, 159.2, 150.5, 141.1, 138.9, 132.5, 130.1, 128.9, 128.4, 128.2, 126.8, 125.3, 125.0, 120.4, 113.6, 52.4, 40.2. \text{ HRMS} (\text{ESI}): \text{ calcd for } \text{C}_{23}\text{H}_{20}\text{N}_3 \text{ [M+H]}^+, 338.1652; \text{ found, } 338.1647.$

N-(2,3-Dihydro-1*H***-inden-2-yl)-2-(pyridin-4-yl)quinazolin-4-amine (11e)**—Brown solid (59 mg, 35%); mp 246–247°C. ¹H NMR (500 MHz, CDCl₃) δ 8.74 (d, *J* = 5.5 Hz, 2H), 8.40 (d, *J* = 5.9 Hz, 2H), 7.93 (d, *J* = 8.2 Hz, 1H), 7.78 – 7.70 (m, 1H), 7.67 (d, *J* = 8.1 Hz, 1H), 7.48 – 7.39 (m, 1H), 7.29 (dd, *J* = 5.2, 3.4 Hz, 2H), 7.22 (dd, *J* = 5.4, 3.3 Hz, 2H), 5.99 (d, *J* = 6.7 Hz, 1H), 5.37 – 5.29 (m, 1H), 3.59 (dd, *J* = 16.2, 7.2 Hz, 2H), 3.07 (dd, *J* = 16.1, 4.8 Hz, 2H). ¹³C NMR (125 MHz, CDCl₃) δ 159.4, 158.4, 150.2, 150.2, 146.4, 141.0, 132.8, 129.2, 126.9, 126.3, 125.0, 122.3, 120.6, 114.1, 77.2, 77.0, 76.7, 52.5, 40.1. HRMS (ESI): calcd for C₂₂H₁₉N₄ [M+H]⁺, 339.1604; found, 339.1608.

N-(2,3-Dihydro-1*H*-inden-2-yl)-2-(thiophen-2-yl)quinazolin-4-amine (11f)—Offwhite solid (68 mg, 40%); mp 243–246°C. ¹H NMR (500 MHz, CDCl₃) δ 8.08 (d, *J* = 3.1 Hz, 1H), 7.85 (d, *J* = 8.4 Hz, 1H), 7.69 (t, *J* = 7.6 Hz, 1H), 7.61 (d, *J* = 8.1 Hz, 1H), 7.45 (d, *J* = 4.9 Hz, 1H), 7.38 – 7.33 (m, 1H), 7.32 – 7.27 (m, 2H), 7.23 (dd, *J* = 5.3, 3.3 Hz, 2H), 7.18 – 7.13 (m, 1H), 5.88 (d, *J* = 6.3 Hz, 1H), 5.31 – 5.24 (m, 1H), 3.60 (dd, *J* = 16.2, 7.2 Hz, 2H), 3.06 (dd, *J* = 16.2, 5.0 Hz, 2H). ¹³C NMR (125 MHz, CDCl₃) δ 159.0, 157.3, 150.4, 145.2, 141.1, 132.6, 129.0, 128.5, 128.2, 127.9, 126.8, 125.1, 124.9, 120.5, 113.5, 52.5, 40.1. HRMS (ESI): calcd for C₂₁H₁₈N₃S [M+H]⁺, 344.1216; found, 344.1218.

N-(2,3-Dihydro-1H-inden-2-yl)-2-(thiophen-3-yl)quinazolin-4-amine (11g)—

Yellow solid (75 mg, 44%); mp 232–233°C. ¹H NMR (500 MHz, CDCl₃) δ 8.34 (d, *J* = 3.0 Hz, 1H), 8.02 (dd, *J* = 5.0, 0.9 Hz, 1H), 7.87 (d, *J* = 8.3 Hz, 1H), 7.74 – 7.65 (m, 1H), 7.62 (d, *J* = 8.0 Hz, 1H), 7.42 – 7.34 (m, 2H), 7.30 (dd, *J* = 5.2, 3.4 Hz, 2H), 7.25 – 7.21 (m, 2H), 5.85 (d, *J* = 6.5 Hz, 1H), 5.36 – 5.25 (m, 1H), 3.58 (dd, *J* = 16.2, 7.2 Hz, 2H), 3.07 (dd, *J* = 16.1, 5.0 Hz, 2H). ¹³C NMR (125 MHz, CDCl₃) δ 159.1, 157.8, 150.5, 143.1, 141.1, 132.5, 128.6, 127.9, 127.4, 126.8, 125.4, 125.1, 124.9, 120.5, 113.5, 52.3, 40.2. HRMS (ESI): calcd for C₂₁H₁₈N₃S [M+H]⁺, 344.1216; found, 344.1215.

N-(2,3-Dihydro-1H-inden-2-yl)-2-(furan-2-yl)quinazolin-4-amine (11h)-Pale-

yellow solid (80 mg, 49%); mp 231–232°C.¹H NMR (500 MHz, CDCl₃) & 7.96 (d, J= 8.4 Hz, 1H), 7.73 – 7.67 (m, 1H), 7.67 – 7.61 (m, 2H), 7.41 – 7.32 (m, 2H), 7.32 – 7.26 (m, 2H), 7.25 – 7.21 (m, 2H), 6.57 (dd, J= 3.3, 1.7 Hz, 1H), 5.98 (d, J= 6.8 Hz, 1H), 5.34 – 5.20 (m, 1H), 3.57 (dd, J= 16.1, 7.2 Hz, 2H), 3.05 (dd, J= 16.1, 5.0 Hz, 2H). ¹³C NMR (125 MHz, CDCl₃) & 159.1, 153.7, 153.4, 150.1, 144.6, 141.0, 132.7, 128.8, 126.8, 125.3, 124.9, 120.5, 113.7, 113.1, 111.8, 52.3, 40.1. HRMS (ESI): calcd for C₂₁H₁₈N₃O [M+H]⁺, 328.1444; found, 328.1448.

Enzymatic Assays

4-Methylumbelliferyl β-D-glucopyranoside (4MU-β-Glc), 4- methylumbelliferyl α-Dglucopyranoside,4-methylumbelliferyl α-D- galactopyranoside, and buffer components were purchased from Sigma-Aldrich (St. Louis, MO). The recombinant wild-type GCase enzyme velaglucerase alfa (Vpriv[®], Shire Human Genetic Therapies, Inc.), acid α-glucosidase enzyme alglucosidase alfa (Lumizyme[®], Genzyme Corporation), α-galactosidase A enzyme agalsidase beta (Fabrazyme[®], Genzyme Corporation) were used in activity assays. The GCase activity assay buffer was composed of 50 mM citric acid, 176 mM K₂HPO₄, and 0.01% Tween-20 at pH5.0, pH 5.9 and pH7.0. A solution of 1 M sodium hydroxide and 1 M glycine (pH 10) was used as the stop solution for all three enzyme activity assays.

GCase Enzyme Activity Assay

The compounds in DMSO solution (0.5 μ L/well) were transferred to a black 96-well plate (the final titration started from 100 μ M, a 12 or 24-point 2-fold dilution series). Enzyme solution (33.5 μ L, 7.5 nM final concentration, in pH 5.9 buffer) was transferred to the wells. After 5 min of incubation at room temperature, the enzyme reaction was initiated by the addition of blue substrate (4MU- β -Glc) (33 μ L/well). The final concentration of the blue substrate was 1.5 mM. The blue substrate reaction was terminated by the addition of 33 μ L/ well stop solution (1 M NaOH and 1 M glycine mixture, pH 10) after 30 min of incubation at 37 °C. The fluorescence was then measured in a Biotek Synergy H1 multi-mode plate reader with Ex = 365 nm and Em = 440 nm. The selected compounds were further assayed under pH 5.0 and pH 7.0 to evaluate their selectivity under various pH conditions.

Enzyme Kinetic Assay²⁰—The substrate resorufin β -D-glucopyranoside was diluted to five concentrations, ranging from 30 to 150 μ M. Seven concentrations of inhibitors (between 0.5- and 5-fold of IC₅₀ value) and a DMSO control were added to the enzyme solution. The final enzyme concentration was 10 nM to give a linear reaction over 10 min. Enzyme kinetics were measured by the addition of 66 μ L of substrate to a 96-well assay plate, followed by 33 μ L of enzyme solution (with or without inhibitor) using a dispense module on a Biotek Synergy H1 multi-mode plate reader. The increase in product fluorescence was measured at 1 min intervals for 10 min in the plate reader. The rate of product formation was calculated by converting the fluorescence units to nanomoles of product per minute using a standard curve of the free fluorophore, resorufin.

Enzyme Selectivity Assays

The acid α -glucosidase and α -galactosidase A enzyme activity assay methods were similar to the GCase enzyme activity assay above with slight modifications. The buffer for the two enzyme assays consisted of 50 mM citric acid, 176 mM K₂HPO₄, and 0.01% Tween-20 at pH 4.8. The final enzyme concentrations for acid α -glucosidase and α -galactosidase A were 8 and 1 nM, respectively. The substrate concentrations for these related enzymes were at 0.16 and 0.4 mM, respectively.

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A robotic pipeline in the High Throughput Analysis Laboratory (HTAL) was used for protein ligand screening by fluorescence thermal shift (FTS) analysis. The pipeline used a Mosquito robot (TTP Labtech) for protein dispensing and an Echo 550 (Labcyte) to add compounds. Thermal scanning coupled with fluorescence detection was performed on a real-time PCR machine CFX384 (Bio-Rad Laboratories). The assay was run in 384-well PCR plates, using 10 μ L citric acid/ K₂HPO₄ buffer (50 mM citric acid, 150 mM K₂HPO₄, pH 5.0) per well. The assay concentration for protein was 1 μ M and that for Sypro Orange (Invitrogen) was 5X. Protein was premixed with Sypro Orange and dispensed to a plate first, and compounds were added. Final concentrations of compounds ranged from 0.5 to 200 μ M. Then the plate was sealed with an optical seal, shaken, and centrifuged. The thermal scan was from 10 to 95 °C with a temperature ramp rate of 1.5 °C/min. The fluorescence was recorded every 10 sec. Data analysis and report generation were performed using the inhouse software excelFTS. The T_m of wild-type GCase was found to follow a logarithmic dose-dependent trend when denaturation was performed in the presence of isofagomine or selected compounds.

N370S Cell Culture and Compound Treatment

The N370S fibroblast cell line was obtained from Coriell, GM00372, cultured in DMEM medium (Life Tech) including 1% v/v L-glutamine 200 mM (Life Tech), 1% v/v pen strep (Life Tech), 10% FBS (Life Tech) at 37 °C and 5% CO₂ and treated with different compounds at indicated concentrations. The same volume of DMSO (0.1% v/v) was used as a control. After a 3-day treatment, cells were washed with inhibitor free media three times and followed by 1% Triton X-100 lysis buffer to lyse cells. Protein concentrations were measured with a Bradford kit (Thermo), and the GCase activity was determined at pH 5.5.

Deglycosylation of Proteins/Molecular Shift Assay ³³

To study the subcellular localization and transport of the various GC mutants (ER and post-ER localization), EndoH and PNGaseF digestions were performed. For both reactions 20 µg of protein was used, and the experimental procedure was performed according to the manufacturer's handbook (New England Biolabs). A positive digestion resulted in a shift in molecular size after the protein was subjected to SDS/PAGE. Anti-hGC (From Johannes Aerts, Leiden University, Leiden, The Netherlands) was used to detect the different forms of GC.

Western Blot

Proteins were denatured in 20% SDS sample buffer at 100 °C for 10 min; 10% Bis-Tris gel (Life Tech) was used for gels; Trans-Blot Turbo PVDF kit (Bio-Rad) was used for membrane transfer. GCase antibody (Sigma-Aldrich) and GAPDH primary antibodies (EMD Millipore) were incubated with the membranes overnight, then the incubated membranes were treated with the secondary antibody (Peroxidase-AffiniPure Goat Anti-Rabbit/Mouse IgG (H+L), Jackson Immunoresearch Lab) for 30 min. Chemidoc MP system (Bio-Rad) was used to scan the membranes and analyze the imaging.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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ABBREVIATIONS USED

GD	Gaucher's disease		
GCase	β-glucocerebrosidase		
ER	endoplasmic reticulum		
ERT'	enzyme replacement therapy		
SRT	substrate reduction therapy		
PC	pharmacological chaperone		
IFG	Isofagomine		
PD	Parkinson's Disease		
DLB	dementia with Lewy bodies		
SAR	structure activity relationship		
GAA	acid a-glucosidase		
GLA	a-galactosidase A		

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Figure 3.

Fluorescent thermal shift analysis of selected compounds. Compound **9a**, **9b**, **11d**, **11f**, **11g**, and IFG showed their ability to stabilize wild-type GCase in a dose-dependent manner. Data represent the results of three independent experiments performed with three replicates per sample.

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Figure 4.

Selectivity of inhibitor **11g** with related hydrolases. **11g** was tested on GCase, acid α -glucosidase (GAA), and α -galactosidase A (GLA). Data represent the results of three independent experiments performed with three replicates per sample.

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B.



Figure 5.

Lineweaver–Burk plots of the enzyme kinetics of GCase inhibitors (A) **4** and (B) **11g**. Each inhibitor was tested in triplicate in two independent assays at the concentrations shown in the legend box of each graph, with (O) indicating the absence of inhibitor. (A) Compounds **4** and (B) Compound **11g** showed an increase in K_m and a decrease in V_{max} , indicating linear mixed inhibition.

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Figure 6.

Modulator **11g** increases GCase protein levels in patient-derived Gaucher's disease fibroblasts. Top: Western blot of N370S fibroblast treated with **11g** or IFG for 3 days. Middle: Western blot of endoH digestion of N370S fibroblast after a 3-day treatment of **11g**, IFG, or DMSO; endoH resistance of proteins indicates their post-ER localization; Bottom: Western blot of PNGase F digestion of N370S fibroblast after a 3-day treatment of **11g**, IFG, or DMSO. The GAPDH signal was used for the loading control, n = 3.

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Scheme 1.

Synthesis of **4**, **6a-6i**, **7a-7i**, **8a-8g**, and **9a-9f** with substituents on the secondary amine Reagents and conditions: (a) (i) sulfolane (ii) PCl₅; (b) RNH₂, K₂CO₃, DMF

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Scheme 2.

Synthesis of **11a-11h** with modifications at the 2-position of the quinazoline ring Reagents and conditions: (a) 2,3-dihydro-1*H*-inden-2-amine, K₂CO₃, DMF; (b) RB(OH)₂, Pd(PPh₃)₄, K₂CO₃, 1,4-dioxane, H₂O

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Table 1

Structure and inhibitory activity of 2-(pyridin-3-yl)quinazoline derivatives with substituted cyclohexyl and related rings^a

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Structure and inhibitory activity of 2-(pyridin-3-yl)quinazoline derivatives with saturated alkyl rings^a







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Table 6

GCase inhibitory activity of 9a, 9b, 9c, 11d, 11g, and 11f at pH 5.0, pH 5.9, and pH 7.0^a

~		IC ₅₀ (nM)		
Comp	Structure	рН 5.0	pH 5.9	pH 7.0
9a		22.4 ± 3.4	8.7 ± 1.1	16.9 ± 2.7
9b		30.1 ± 4.5	9.9 ± 1.3	18.8 ± 2.5
9c		25.9 ± 3.9	8.3 ± 1.0	12.7 ± 1.4
11d		10.5 ± 1.5	6.5 ± 0.7	9.4 ± 1.4
11f	N N HN K	13.9 ± 1.9	8.2 ± 1.0	12.9 ± 2.0
11g		8.9 ± 1.3	5.2 ± 0.6	6.6 ± 1.0

^{*a*}Experiments were performed in triplicate, and the mean \pm SD is shown.