



Discovery of 4-amino-1*H*-pyrazolo[3,4-*d*]pyrimidin derivatives as novel discoidin domain receptor 1 (DDR1) inhibitors

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ABSTRACT

DDR1 is a receptor tyrosine kinase that is activated by triple-helical collagens and has become an attractive target for anticancer therapy given its involvement in tumor growth, metastasis development, and tumor dormancy. Several drugs on the market, such as dasatinib and nilotinib, were reported to potently suppress the function of DDR1 and show significant therapeutic benefits in a variety of preclinical tumor models. Whereas only a few selective DDR1 inhibitors were disclosed in recent years. A series of 4-amino-1*H*-pyrazolo[3,4-*d*]pyrimidin derivatives were designed and synthesized. All compounds were evaluated via DDR1 kinase inhibition assay and cell anti-proliferative assay. One of the representative compounds, **6c**, suppressed DDR1 kinase activity with an IC₅₀ value of 44 nM and potently inhibited cell proliferation in DDR1-overexpressing cell lines HCT-116 and MDA-MB-231 with IC₅₀ value of 4.00 and 3.36 μM respectively. Further molecular docking study revealed that **6c** fitted ideally into DDR1 binding pocket and maintained the crucial hydrogen bonds with DDR1 kinase domain. Overall, these results suggest that the compound **6c** is a potential DDR1 inhibitor deserving further investigation for cancer treatment.

1. Introduction

Discoidin domain receptors (DDRs), classified into two types: DDR1 and DDR2, are receptor tyrosine kinases (RTKs)¹. DDrs regulate basic aspects of cell behaviors (cellular morphogenesis, differentiation, proliferation, adhesion, migration, and invasion) by interacting with collagen and cross-talking with different types of cell surface receptors (such as integrins and other RTKs)². Increasing studies have reported that DDrs play the part of potential oncogenic drivers in various cancers, including lung cancer, breast cancer, brain cancer, prostate cancer, leukemia, colorectal cancer, liver cancer, and so on³. Moreover, DDrs are linked closely with fibrotic disorders⁴⁻⁶ and atherosclerosis⁷⁻⁹. A survey revealed that nilotinib strongly inhibited human colon cancer cells (CRC) invasion in vitro and reduced their metastatic potential in intrasplenic tumor mouse models and patient-derived metastatic and circulating CRC cell lines by inhibiting the kinase activity of DDR1¹⁰. Other surveys showed that inhibition of DDR1 sensitized glioblastoma cells to radio- and chemotherapy by inducing autophagy^{11,12}. Silencing DDR1 by siRNA reduced metastatic activity in lung cancer models and enhanced the chemosensitivity of breast cancer cells to genotoxic drugs

¹³⁻¹⁵. Therefore, DDR1 has been considered as a promising drug target for anticancer or other diseases' therapy.

Until recently, several small molecules¹⁶⁻²¹ were reported to potently and selectively suppress the kinase activity of DDR1 (Fig. 1), with IC₅₀ of low nanomole. In 2013, the compound **7rh** as a first selective DDR1 inhibitor was reported to effectively inhibit the proliferation, invasion, adhesion, and tumorigenicity of cancer cells with high DDR1 expression¹⁶. A research showed **7rh** strongly inhibited gastric cancer (GC) tumor growth, supporting the potential of DDR1-induced mechanisms for clinical GC therapy²². The compound **5** potently suppressed collagen-induced DDR1 signaling and epithelial-mesenchymal transition, and exhibited promising therapeutic efficacy in orthotopic mouse models of pancreatic cancer²¹. All those surveys show that DDR1 is becoming an attractive target for drug discovery.

There will still be a rise in the demand for DDR1 targeted therapies. A series of 4-amino-1*H*-pyrazolo[3,4-*d*]pyrimidin derivatives were designed and synthesized (Scheme 1). Based on high homology in the ATP-binding pocket, core scaffolds of tyrosine kinase inhibitors were screened to occupy the adenine portion of the DDR1 ATP-binding pocket using Discovery Studio 3.0, and new inhibitors were eventually designed

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by combining the core scaffold screened and the fragment of DDR1-IN-1 (Fig. 2). Moreover, the potential efficacy of these compounds was evaluated in vitro.

2. Results and discussions

2.1. Chemistry

The general schemes for the synthesis of compounds **6a**–**6u** were summarized in Scheme 1. The key steps were copper-catalyzed Ullmann coupling reaction²³ and palladium-catalyzed Suzuki coupling reaction²⁴. Briefly, commercially available 2-iodo-4-nitro-toluene (**7**) was treated with phenol (**8**) under copper catalysis to afford the Ullmann coupling product (**9**), which went through iodization to yield the compound (**10**). Then, the intermediate (**10**) was reacted with bis(pinacolato)diboron under palladium catalysis to produce the compound (**11**), which was hydrolyzed to obtain the product (**12**). The compound (**12**) went through palladium-catalyzed Suzuki coupling reaction with commercially available compound (**13**) to afford the intermediate (**14**), which was reduced to the aromatic amine (**15**). In the end, the intermediate (**15**) and the synthesized acyl chloride (**16**) went through amide condensation to afford the target compounds (**6a**–**6u**).

2.2. In vitro kinase inhibitory activities

To explore the inhibitory activities of the designed compounds, these compounds were evaluated using the LANCE ULTRA kinase assay. The results are summarized in Table 1. **GZD856** was used as a positive control for its strong binding affinity to DDR1²⁵. Compounds **6c**, **6d**, and **6u** inhibited DDR1 with IC₅₀ values ranging from 0.044 to 0.070 μM. Accordingly, substituents of hydrogen (**6u**) or methyl (**6c**) group at R₁ could be tolerated without reducing the inhibitory activity against DDR1. Whereas it seemed that substituents on the *N*-terminal phenyl ring dramatically impacted DDR1 kinase inhibition. When (4-methylpiperazin-1-yl)methylene substituent (**6c**, IC₅₀ = 44 nM) was replaced by (4-ethylpiperazin-1-yl)methylene substituent (**6d**, IC₅₀ = 70 nM) at R₃, the DDR1 inhibitory activity was reduced. When (4-methylpiperazin-1-yl)methylene group was introduced at R₂, the resulting compound (**6e**) showed twice as potent as the compound (**6a**) containing (4-methylpiperazin-1-yl)methylene group at R₃. However, the potency of compounds (**6j**, **6k**, **6l**, and **6m**) which contained other alicyclic amines at R₃ was significantly decreased. When iodo group was introduced at R₂, the IC₅₀ value of the compound (**6i**) was 131.0 nM, less potent than **6c**. Besides, the compounds (**6p**, **6r**, and **6t**) with halo substituents at R₂ displayed significantly weaker DDR1 inhibitory activities than the compound (**6k**) with trifluoromethyl substituent. As a result, the compound with trifluoromethyl substituent at R₂ and (4-methylpiperazin-1-yl)methylene group at R₃ showed the most potent DDR1 inhibitory activity.

2.3. Inhibition of cell proliferation

To determine their inhibitory activities of cell proliferation, all designed compounds were evaluated against two cancer cell lines expressing high level of DDR1 by MTT assay. The results are shown in Table 2, and **nilotinib** was selected as the positive control because of its strong DDR1 inhibitory activity. As shown in Table 2, all of the compounds were evaluated their inhibition of DDR1-overexpressing cancer cell lines and most of them exhibited significantly anti-proliferative activities (>95%) at the concentration of 10 μM. Then the compounds, inhibiting >95% cell proliferation, were tested their IC₅₀ values. It was surprising that most of the compounds showed better anti-proliferative activities than **nilotinib**. Especially, compound **6c** potently inhibited cell proliferation in cell lines HCT-116 and MDA-MB-231 with IC₅₀ values of 4.00 and 3.36 μM respectively. The results indicated that **6c** could be a promising lead compound for cancer treatment.

2.4. Molecular docking study

To further elucidate the interaction between the compound **6c** and DDR1, Discovery Studio 4.0/CDOCKER protocol was employed to analyze the possible binding mode. The binding mode of the most potent compound **6c** and DDR1 is shown in Fig. 3. The results suggested that the compound **6c** could bind tightly with the DDR1 kinase via four hydrogen bonds, two π-π stacked interactions, electrostatic interactions, and many Van der Waals interactions. As expected, the nitrogen atom and the secondary amino hydrogen atom at the 1*H*-pyrazolo[3,4-*d*]pyrimidin moiety formed two essential hydrogen bonds with the secondary amino hydrogen atom (2.4 Å, 173.1°) and the carbonyl oxygen atom (2.3 Å, 134.7°) of Met⁷⁰⁴ in the hinge binding area respectively, which might play a deciding role in the binding affinity with DDR1 kinase. Besides, 1*H*-pyrazolo[3,4-*d*]pyrimidin generated π-π interactions with the phenyl ring of the residue Tyr⁷⁰³ (Figure C), indicating that the 1*H*-pyrazolo[3,4-*d*]pyrimidin moiety plays an important role in the combination of the receptor and the ligand. Moreover, Glu⁶⁷² in the αC-helix and Asp⁷⁸⁴ in the DFG motif formed two hydrogen bonds with the linker amide moiety, the distances of which were 2.3 Å (127.5°) and 2.1 Å (170.4°) respectively, contributing to the affinity increment of compound **6c** with DDR1. These results revealed that the compound **6c** had a good binding ability to DDR1 kinase as designed and explained the high potency of inhibition on DDR1 Kinase.

3. Conclusion

In summary, a new class of 4-amino-1*H*-pyrazolo[3,4-*d*]pyrimidin compounds have been designed and synthesized, and their DDR1 inhibitory activities and anti-proliferative activities were evaluated in vitro. It was found that compound **6c** exhibited robust potency of DDR1 inhibition with an IC₅₀ value of 44 nM. Besides, compound **6c** potently

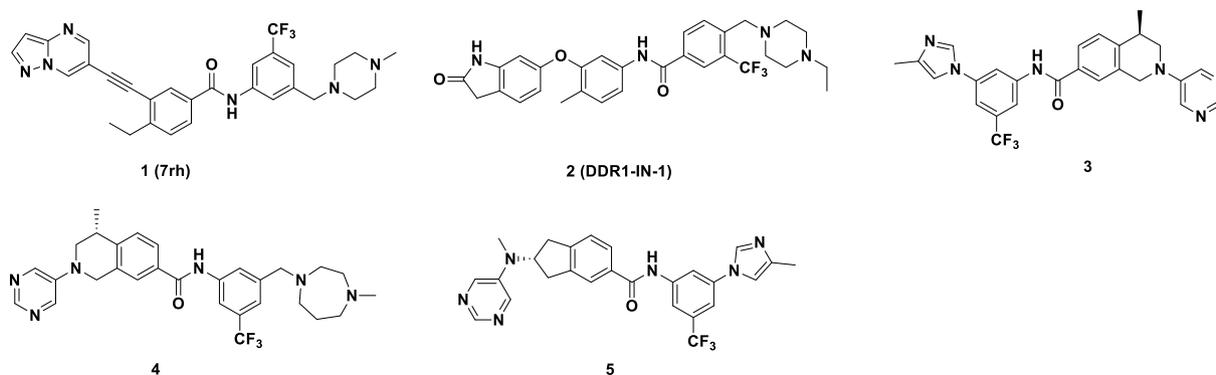


Fig. 1. Part of reported selective DDR1 inhibitors.

inhibited proliferation of cancer cell lines expressing high levels of DDR1 with IC₅₀ values in low micromole range. These findings demonstrate that the compound **6c** can be a potent DDR1 inhibitor and worthy of further investigation.

4. Experiment

4.1. Chemistry

All reagents and solvents were purchased from commercial sources without further purification. Flash chromatography was performed using 300-mesh silica gel. All reactions were monitored by TLC using silica gel plates with fluorescence F254 and UV-light visualization. ¹H NMR spectra and ¹³C NMR spectra were recorded on a Bruker AV-300 spectrometer at 300 MHz. Coupling constants (*J*) are expressed in hertz (Hz). Chemical shifts (δ) of NMR are reported in parts per million (ppm) relative to an internal standard (TMS). ESI-MS spectra were recorded on an Agilent/HP 1100 Series LC/MSD Trap SL mass spectrometer. High resolution mass spectrometry (HRMS) was obtained on a Q-tof high resolution mass spectrometer. Melting points (uncorrected) were determined on a RY-1 MP apparatus. The IR spectra was recorded on a Nicolet Impact 410 infrared spectrometer.

4.2. General procedure for the preparation of the compounds (**6a**–**6u**)

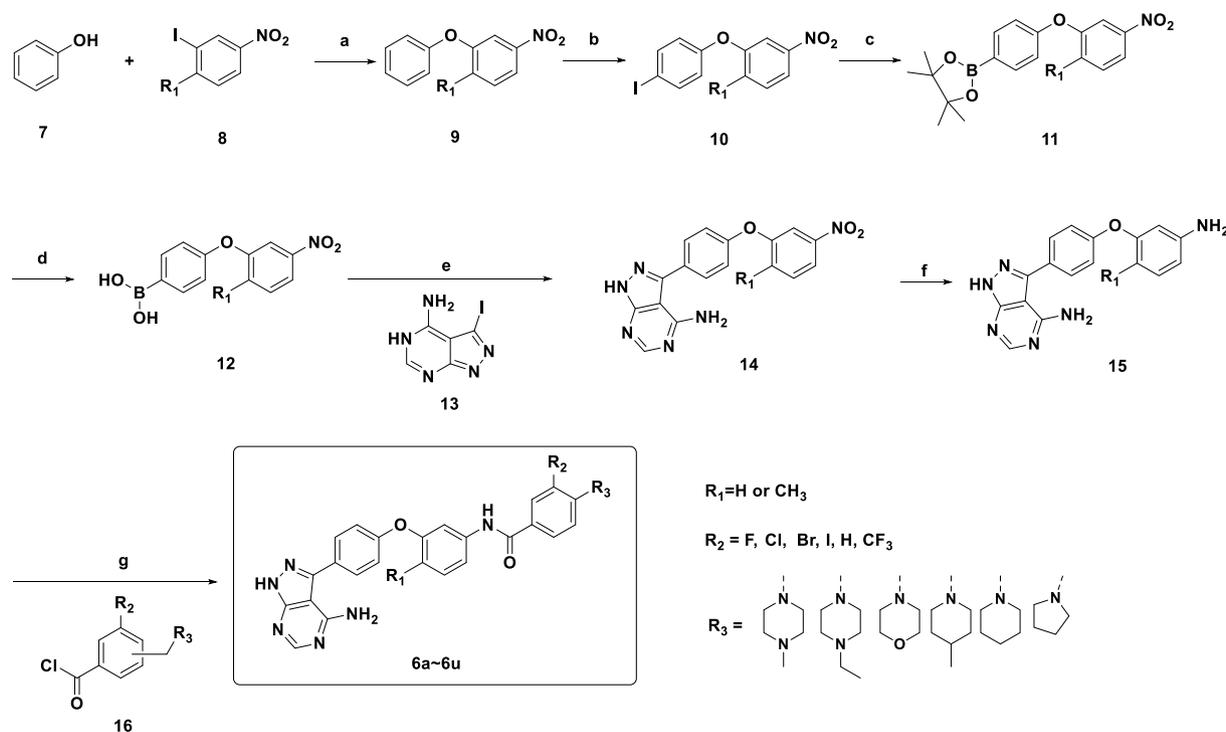
1-methyl-4-nitro-2-phenoxybenzene (9). A mixture of 2-iodo-4-nitrotoluene (**7**) (1052.1 mg, 4 mmol), phenol (**8**) (564.7 mg, 6 mmol), Cs₂CO₃ (2606.6 mg, 8 mmol), CuI (152.4 mg, 0.8 mmol), and *N,N*-dimethylglycine hydrochloride (355.0 mg, 2.4 mmol), and anhydrous 1,4-dioxane (15 mL) in a schlink tube was heated at 135 °C under nitrogen atmosphere. After the reaction completed monitored by TLC, the cooled mixture was partitioned between ethyl acetate and water. The organic layer was separated, and the aqueous layer was extracted with ethyl acetate. The combined organic layers were washed with brine, dried over anhydrous Na₂SO₄, and concentrated in vacuo. The residual

oil was loaded on a silica gel column and eluted with ethyl acetate/petroleum ether (1/100) to afford the product (778.9 mg, 85.0%). ¹H NMR (300 MHz, Chloroform-*d*) δ 7.95 (dd, *J* = 8.3, 2.3 Hz, 1H), 7.70 (d, *J* = 2.3 Hz, 1H), 7.48–7.40 (m, 3H), 7.23 (tt, *J* = 7.2, 1.0 Hz, 1H), 7.07–7.01 (m, 2H), 2.45 (s, 3H).

2-(4-iodophenoxy)-1-methyl-4-nitrobenzene (10). To a 100 mL round-bottomed flask were added 1-methyl-4-nitro-2-phenoxybenzene (1507.9 mg, 6.58 mmol), *N*-iodosuccinimide (1481.0 g, 6.71 mmol), FeCl₃ (107.1 mg, 0.66 mmol), and acetonitrile (20 mL), and the mixture was stirred at 60 °C. After the reaction completed monitored by TLC, ethyl acetate was added, and the organic phase was washed with brine (3 times), which was dried over anhydrous Na₂SO₄. The residual oil was loaded on a silica gel column and eluted with ethyl acetate/petroleum ether (1/100) to afford the product (1844.4 mg, 80.0%). ¹H NMR (300 MHz, Chloroform-*d*) δ 7.94 (dd, *J* = 8.3, 2.0 Hz, 1H), 7.74–7.63 (m, 3H), 7.42 (d, *J* = 8.3 Hz, 1H), 6.77 (d, *J* = 8.8 Hz, 2H), 2.39 (s, 3H).

4,4,5,5-tetramethyl-2-(4-(2-methyl-5-nitrophenoxy)phenyl)-1,3,2-dioxaborolane (11). To a solution of 2-(4-iodophenoxy)-1-methyl-4-nitrobenzene (1844.4 mg, 5.20 mmol) in anhydrous 1,4-dioxane (20 mL) were added 1,1'-bis(diphenylphosphino)ferrocene dichloropalladium (II) (**II**) (380.5 mg, 0.1 mmol), bis(pinacolato)diboron (1979.2 mg, 1.5 mmol), and potassium acetate (1531.0 mg, 15.6 mmol). The reaction mixture was stirred at 101 °C. After the reaction completed monitored by TLC, the cooled mixture was filtered. The filtrate was concentrated in vacuo, resuspended in ethyl acetate and water, and separated. The organic layer was extracted with brine and dried over anhydrous Na₂SO₄. Purification by silica gel chromatography with ethyl acetate/petroleum ether (1/20) provided the product as white solid (1613.0 mg, 87.3%). ¹H NMR (300 MHz, Chloroform-*d*) δ 7.92 (dd, *J* = 8.4, 2.3 Hz, 1H), 7.86–7.79 (m, 2H), 7.69 (d, *J* = 2.3 Hz, 1H), 7.40 (d, *J* = 8.4 Hz, 1H), 6.99–6.92 (m, 2H), 2.36 (s, 3H), 1.35 (s, 12H).

4-(2-methyl-5-nitrophenoxy)phenylboronic acid (12). 4,4,5,5-tetramethyl-2-(4-(2-methyl-5-nitrophenoxy)phenyl)-1,3,2-dioxaborolane (441.4 mg, 1.24 mmol) and sodium periodate (795.7 mg, 3.72 mmol) were stirred in 5 mL of a 4:1 mixture of THF and water for 30 min, at



Scheme 1. Synthesis of 4-amino-1H-pyrazolo[3,4-d]pyrimidin derivatives. Reagents and conditions: (a) Cs₂CO₃, CuI, *N,N*-dimethylglycine hydrochloride, Dioxane, 135 °C; (b) NIS, FeCl₃, CH₃CN, 60 °C; (c) Pd(dppf)Cl₂, KOAc, Dioxane, 100 °C; (d) NaIO₄, 1 M HCl, THF: H₂O (4: 1), 25 °C; (e) Pd(dppf)Cl₂, K₃PO₄·3H₂O, THF/H₂O = 4/1, 100 °C; (f) activated carbon, FeCl₃, 80% Hydrazine Hydrate, EtOH, 85 °C; (g) TEA, absolute THF, 65 °C

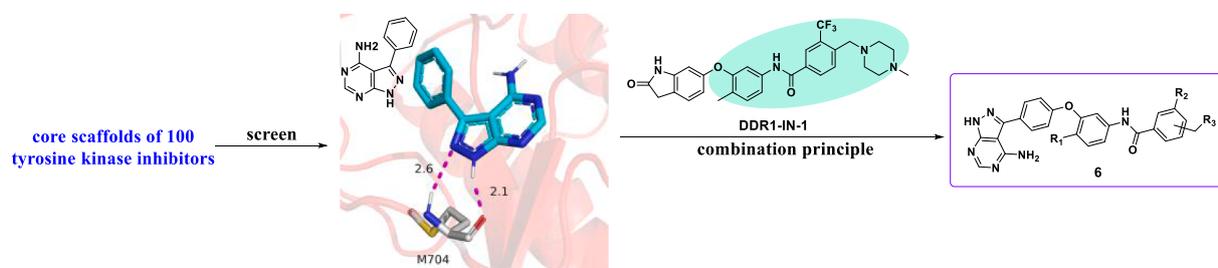


Fig. 2. Design of 4-amino-1H-pyrazolo[3,4-d]pyrimidin derivatives as novel DDR1 inhibitors.

Table 1

In vitro kinase inhibitory activities of compounds 6a~6u against DDR1.

Cpds	R ₁	R ₂	R ₃	IC ₅₀ (μM)	Cpds	R ₁	R ₂	R ₃	IC ₅₀ (μM)
6a	CH ₃	H		2.694	6l	CH ₃	CF ₃		0.975
6b	CH ₃	H		N.D.	6m	CH ₃	CF ₃		2.810
6c	CH ₃	CF ₃		0.044	6n	CH ₃	H		N.D.
6d	CH ₃	CF ₃		0.070	6o	CH ₃	F		N.D.
6e	CH ₃		H	1.213	6p	CH ₃	F		24.350
6f	CH ₃	H		N.D.	6q	CH ₃	Cl		N.D.
6g	CH ₃	H		N.D.	6r	CH ₃	Cl		2.172
6h	CH ₃	H		N.D.	6s	CH ₃	Br		N.D.
6i	CH ₃	I		0.131	6t	CH ₃	Br		2.468
6j	CH ₃	CF ₃		1.189	6u	H	CF ₃		0.044
6k	CH ₃	CF ₃		0.416	GZD856				0.005

DDR1 experiments were performed using the LANCE ULTRA kinase assay, according to the manufacturer's instructions. The data is from one experiment. N.D. = Not determined.

which time aqueous hydrochloric acid (1 N, 0.87 mL, 0.87 mmol) was added to the suspension. The reaction mixture was stirred at ambient temperature. After TLC showed complete consumption of the arylboronic acid ester, the reaction mixture was diluted with water and extracted with ethyl acetate. The combined extracts were washed with water and brine, dried over sodium sulfate, filtered, and concentrated to dryness by rotary evaporation. The residue was purified by silica gel chromatography with ethyl acetate/petroleum ether (1/2), giving the product as white solid (185.6 mg, 54.8%).

3-(4-(2-methyl-5-nitrophenoxy)phenyl)-1H-pyrazolo[3,4-d]pyrimidin-4-amine (14). To a mixture of 3-iodo-5H-pyrazolo[3,4-d]pyrimidin-4-amine (871.6 mg, 3.34 mmol) and THF/H₂O (40 mL, v/v, 4/1) were added 4-(2-methyl-5-nitrophenoxy)phenylboronic acid (1184.4 mg, 4.34 mmol), K₃PO₄·3H₂O (2663.1 mg, 10.0 mmol), and Pd(dppf)Cl₂ (490.2 mg, 0.67 mmol). The reaction mixture was placed into an oil bath preheated to 100 °C, with stirring at this temperature under nitrogen. After the reaction completed monitored by TLC, the resulting mixture was filtered through a celite pad and concentrated to dryness by rotary evaporation. The residue was purified by silica gel chromatography with DCM/MeOH (80/1, 1% TEA) to afford the product as a white color solid (622.3 mg, 51.42%). ¹H NMR (300 MHz, DMSO-*d*₆) δ 13.64 (s, 1H), 8.26 (s, 1H), 8.05 (dd, *J* = 8.4, 2.3 Hz, 1H), 7.87–7.66 (m, 4H), 7.36–7.14 (m,

3H), 2.43 (s, 3H).

3-(4-(5-amino-2-methylphenoxy)phenyl)-1H-pyrazolo[3,4-d]pyrimidin-4-amine (15). The solution of 3-(4-(2-methyl-5-nitrophenoxy)phenyl)-1H-pyrazolo[3,4-d]pyrimidin-4-amine (622.3 mg, 1.72 mol) and ethanol (26.8 mL) was stirred at 60 °C, an appropriate amount of activated carbon (60.2 mg) and ferric chloride (62.0 mg) were added at the temperature, the mixture was heated to 85 °C and 80 percent hydrazine hydrate (5.7 mL) was added to the solution. The reaction mixture then was refluxed for 2 h and monitored by TLC. The mixture was filtered and the precipitate was washed with ethanol. The filtrate was concentrated under reduced pressure and the residue was poured into water with stirring for 30 min. The precipitate was filtered and dried to furnish the product as white solid (480.3 mg, 84.0%). ¹H NMR (300 MHz, DMSO-*d*₆) δ 13.54 (s, 1H), 8.23 (s, 1H), 7.64 (d, *J* = 6.9 Hz, 2H), 7.13–6.90 (m, 3H), 6.39 (d, *J* = 6.1 Hz, 1H), 6.25 (s, 1H), 5.05 (s, 2H), 2.03 (s, 3H).

N-(3-(4-(4-amino-1H-pyrazolo[3,4-d]pyrimidin-3-yl)phenoxy)-4-methylphenyl)-4-((4-methylpiperazin-1-yl)methyl)benzamide (6a). A 25 mL round bottom flask charged with 3-(4-(5-amino-2-methylphenoxy)phenyl)-1H-pyrazolo[3,4-d]pyrimidin-4-amine (7) (50.0 mg, 0.18 mmol), THF (5 mL), Et₃N (0.05 mL, 0.36 mmol) treated with 4-((4-methylpiperazin-1-yl)methyl)benzoyl chloride (8) (about 0.25 mmol).

Table 2

The anti-proliferative effects of compounds on two cancer cells harboring high levels of DDR1.

Cpds	HCT-116		MDA-MB-231	
	Inhibition at 10 μ M (%)	IC ₅₀ (μ M)	Inhibition at 10 μ M (%)	IC ₅₀ (μ M)
6a	26.70 \pm 8.01	>10	94.54 \pm 0.57	N.D.
6b	99.59 \pm 0.21	2.00 \pm 0.19	98.70 \pm 0.37	2.68 \pm 0.26
6c	99.96 \pm 0.04	4.00 \pm 0.12	97.93 \pm 1.15	3.36 \pm 0.11
6d	99.74 \pm 0.09	5.48 \pm 0.21	99.28 \pm 0.77	3.85 \pm 0.07
6e	99.83 \pm 0.04	4.47 \pm 0.28	99.28 \pm 0.38	N.D.
6f	99.94 \pm 0.30	1.37 \pm 0.03	97.40 \pm 0.37	2.76 \pm 0.02
6	99.96 \pm 0.11	3.04 \pm 0.04	98.83 \pm 0.55	N.D.
6h	97.25 \pm 0.89	4.89 \pm 0.08	98.96 \pm 0.74	3.53 \pm 0.08
6i	99.74 \pm 0.09	4.51 \pm 0.02	98.87 \pm 0.19	3.21 \pm 0.07
6j	34.86 \pm 0.76	>10	98.47 \pm 0.00	N.D.
6k	20.50 \pm 1.74	>10	71.28 \pm 2.87	N.D.
6l	36.91 \pm 8.32	>10	29.62 \pm 0.96	>10
6m	73.73 \pm 1.16	N.D.	42.06 \pm 2.10	>10
6n	33.63 \pm 1.00	>10	88.67 \pm 1.29	N.D.
6o	73.25 \pm 3.68	N.D.	92.06 \pm 0.18	N.D.
6p	11.09 \pm 7.21	>10	93.06 \pm 0.00	N.D.
6q	99.91 \pm 0.15	2.91 \pm 0.12	99.22 \pm 0.37	3.83 \pm 0.39
6r	58.26 \pm 6.05	N.D.	96.48 \pm 0.92	N.D.
6s	99.55 \pm 0.05	6.56 \pm 0.80	100.00 \pm 0.37	N.D.
6t	35.02 \pm 9.04	>10	81.29 \pm 2.87	N.D.
6u	99.85 \pm 0.16	5.96 \pm 0.06	98.20 \pm 0.38	4.29 \pm 0.01
Nilotinib	88.14 \pm 0.02	5.75 \pm 0.10	93.60 \pm 0.77	7.31 \pm 0.17

Biological activity data on inhibition of cancer cell lines were expressed as the means \pm SD derived from three independent measurements. N.D. = not determined.

The reaction mixture was stirred at room temperature. After the reaction completed monitored by TLC, then Saturated sodium bicarbonate solution was added. The reaction mixture was extracted with ethyl acetate. The organic layers were combined and washed with brine, and then dried over anhydrous Na₂SO₄. The resulting residue was purified by TLC on a silica gel (DCM: MeOH = 40: 1, 2% TEA), then was beaten in CH₃CN at 80 °C for 12 h. The mixture was cooled to room temperature and filtered to afford the product as white solid (17.5 mg, 21.26%). Mp: 179–180 °C. ¹H NMR (300 MHz, DMSO-*d*₆) δ 13.56 (s, 1H), 10.24 (s, 1H), 8.21 (s, 1H), 7.87 (d, *J* = 7.9 Hz, 2H), 7.66 (d, *J* = 8.2 Hz, 2H), 7.57 (dd, *J* = 5.3, 1.7 Hz, 2H), 7.42 (d, *J* = 7.9 Hz, 2H), 7.32 (d, *J* = 8.7 Hz, 1H), 7.08 (d, *J* = 8.4 Hz, 2H), 6.79 (s, 1H), 3.51 (d, *J* = 4.8 Hz, 2H), 2.36 (s, 7H), 2.18 (d, *J* = 8.5 Hz, 6H). HRMS (ESI+) *m/z* calculated for C₃₁H₃₂N₈O₂ [M+H]⁺, 549.2721; found, 549.2723.

N-(3-(4-(4-amino-1H-pyrazolo[3,4-d]pyrimidin-3-yl)phenoxy)-4-methylphenyl)-4-(piperidin-1-yl-methyl)benzamide (6b). Prepared in the same manner as 6a. Yield: 26.8%, mp: 204–205 °C. ¹H NMR (300 MHz, DMSO-*d*₆) δ 13.56 (s, 1H), 10.24 (s, 1H), 8.21 (s, 1H), 7.86 (d, *J* = 7.8 Hz, 2H), 7.66 (d, *J* = 8.5 Hz, 2H), 7.57 (d, *J* = 6.1 Hz, 2H), 7.41 (d, *J* = 7.8 Hz, 2H), 7.32 (d, *J* = 8.6 Hz, 1H), 7.08 (d, *J* = 8.6 Hz, 2H), 3.49 (s, 2H), 2.32 (s, 4H), 2.19 (s, 3H), 1.49 (s, 4H), 1.38 (s, 2H). HRMS (ESI+) *m/z* calculated for C₃₁H₃₁N₇O₂ [M+H]⁺, 534.2612; found, 534.2620.

N-(3-(4-(4-amino-1H-pyrazolo[3,4-d]pyrimidin-3-yl)phenoxy)-4-methylphenyl)-4-((4-methyl-piperazin-1-yl)methyl)-3-(trifluoromethyl)benzamide (6c). Prepared in the same manner as 6a. Yield: 23.2%, mp: 160–161 °C. ¹H NMR (300 MHz, DMSO-*d*₆) δ 13.56 (s, 1H), 10.46 (s, 1H), 8.25–8.13 (m, 3H), 7.90 (d, *J* = 8.0 Hz, 1H), 7.72–7.62 (m, 2H),

7.62–7.50 (m, 2H), 7.34 (d, *J* = 8.2 Hz, 1H), 7.16–7.04 (m, 2H), 6.81 (s, 1H), 3.67 (s, 2H), 2.40 (s, 8H), 2.19 (d, *J* = 7.2 Hz, 6H). HRMS (ESI+) *m/z* calculated for C₃₂H₃₁F₃N₈O₂ [M+H]⁺, 617.2595; found, 617.2590.

N-(3-(4-(4-amino-1H-pyrazolo[3,4-d]pyrimidin-3-yl)phenoxy)-4-methylphenyl)-4-((4-ethyl-piperazin-1-yl)methyl)-3-(trifluoromethyl)benzamide (6d). Prepared in the same manner as 6a. Yield: 22.8%, mp: 158–160 °C. ¹H NMR (300 MHz, DMSO-*d*₆) δ 13.54 (s, 1H), 10.44 (s, 1H), 8.28–8.09 (m, 3H), 7.88 (d, *J* = 8.0 Hz, 1H), 7.65 (d, *J* = 8.3 Hz, 2H), 7.60–7.46 (m, 2H), 7.32 (d, *J* = 8.2 Hz, 1H), 7.07 (d, *J* = 8.2 Hz, 2H), 6.78 (s, 1H), 3.64 (s, 2H), 2.45–2.26 (m, 8H), 2.18 (s, 3H), 0.96 (t, *J* = 7.0 Hz, 3H). HRMS (ESI+) *m/z* calculated for C₃₃H₃₃F₃N₈O₂ [M+H]⁺, 631.2751; found, 631.2752.

N-(3-(4-(4-amino-1H-pyrazolo[3,4-d]pyrimidin-3-yl)phenoxy)-4-methylphenyl)-3-((4-methyl-piperazin-1-yl)methyl)benzamide (6e). Prepared in the same manner as 6a. Yield: 42.0%, mp: 166–168 °C. ¹H NMR (300 MHz, DMSO-*d*₆) δ 13.55 (s, 1H), 10.26 (s, 1H), 8.22 (s, 1H), 7.81 (d, *J* = 6.8 Hz, 2H), 7.67 (d, *J* = 8.2 Hz, 2H), 7.58 (d, *J* = 6.5 Hz, 2H), 7.46 (d, *J* = 8.6 Hz, 2H), 7.33 (d, *J* = 8.5 Hz, 1H), 7.09 (d, *J* = 8.2 Hz, 2H), 6.81 (s, 1H), 3.51 (s, 2H), 2.36 (s, 7H), 2.20 (s, 3H), 2.14 (s, 3H). HRMS (ESI+) *m/z* calculated for C₃₁H₃₂N₈O₂ [M+H]⁺, 549.2721; found, 549.2728.

N-(3-(4-(4-amino-1H-pyrazolo[3,4-d]pyrimidin-3-yl)phenoxy)-4-methylphenyl)-4-((4-methyl-piperidin-1-yl)methyl)benzamide (6f). Prepared in the same manner as 6a. Yield: 13.2%, mp: 198–199 °C. ¹H NMR (300 MHz, DMSO-*d*₆) δ 13.58 (s, 1H), 10.26 (s, 1H), 8.24 (s, 1H), 7.89 (d, *J* = 8.2 Hz, 2H), 7.74–7.65 (m, 2H), 7.61 (dq, *J* = 4.7, 2.1 Hz, 2H), 7.44 (d, *J* = 8.0 Hz, 2H), 7.35 (d, *J* = 8.9 Hz, 1H), 7.16–7.08 (m, 2H), 6.82 (s, 1H), 3.52 (s, 2H), 2.78 (d, *J* = 11.1 Hz, 2H), 2.22 (s, 3H), 1.95 (t, *J* = 11.2 Hz, 2H), 1.58 (dd, *J* = 12.2, 3.3 Hz, 2H), 1.32 (s, 1H), 1.18 (td, *J* = 12.0, 3.7 Hz, 2H), 0.91 (d, *J* = 6.3 Hz, 3H). HRMS (ESI+) *m/z* calculated for C₃₂H₃₃N₇O₂ [M+H]⁺, 548.2768; found, 548.2779.

N-(3-(4-(4-amino-1H-pyrazolo[3,4-d]pyrimidin-3-yl)phenoxy)-4-methylphenyl)-4-(pyrrolidin-1-yl-methyl)benzamide (6g). Prepared in the same manner as 6a. Yield: 43.5%, mp: 166–167 °C. ¹H NMR (300 MHz, DMSO-*d*₆) δ 13.58 (s, 1H), 10.25 (s, 1H), 8.24 (s, 1H), 7.90 (d, *J* = 7.8 Hz, 2H), 7.69 (d, *J* = 8.2 Hz, 2H), 7.61 (s, 2H), 7.45 (d, *J* = 7.8 Hz, 2H), 7.34 (d, *J* = 8.5 Hz, 1H), 7.11 (d, *J* = 8.1 Hz, 2H), 6.78 (s, 1H), 3.66 (s, 2H), 2.46 (s, 4H), 2.22 (s, 3H), 1.72 (s, 4H). HRMS (ESI+) *m/z* calculated for C₃₀H₂₉N₇O₂ [M+H]⁺, 520.2455; found, 520.2460.

N-(3-(4-(4-amino-1H-pyrazolo[3,4-d]pyrimidin-3-yl)phenoxy)-4-methylphenyl)-4-(morpholino-methyl)benzamide (6h). Prepared in the same manner as 6a. Yield: 7.7%, mp: 182–184 °C. ¹H NMR (300 MHz, DMSO-*d*₆) δ 13.59 (s, 1H), 10.30 (s, 1H), 8.25 (s, 1H), 7.94 (d, *J* = 7.7 Hz, 2H), 7.70 (d, *J* = 8.3 Hz, 2H), 7.61 (s, 2H), 7.52 (s, 2H), 7.36 (d, *J* = 8.8 Hz, 1H), 7.12 (d, *J* = 8.3 Hz, 2H), 3.64 (s, 6H), 2.44 (s, 3H), 2.23 (s, 3H). HRMS (ESI+) *m/z* calculated for C₃₀H₂₉N₇O₃ [M+H]⁺, 536.2405; found, 536.2411.

N-(3-(4-(4-amino-1H-pyrazolo[3,4-d]pyrimidin-3-yl)phenoxy)-4-methylphenyl)-3-iodo-4-((4-methyl-piperazin-1-yl)methyl)benzamide (6i). Prepared in the same manner as 6a. Yield: 49.9%, mp: 166–167 °C. ¹H NMR (300 MHz, DMSO-*d*₆) δ 13.57 (s, 1H), 10.33 (s, 1H), 8.37 (s, 1H), 8.22 (s, 1H), 7.91 (d, *J* = 8.0 Hz, 1H), 7.67 (d, *J* = 8.2 Hz, 2H), 7.57 (d, *J* = 8.9 Hz, 2H), 7.50 (d, *J* = 7.8 Hz, 1H), 7.33 (d, *J* = 8.2 Hz, 1H), 7.09 (d, *J* = 8.3 Hz, 2H), 3.50 (s, 2H), 2.44 (s, 4H), 2.35 (s, 4H), 2.18 (d, *J* = 9.4 Hz, 6H). HRMS (ESI+) *m/z* calculated for C₃₁H₃₁I_N₈O₂ [M+H]⁺, 675.1687; found, 675.1689.

N-(3-(4-(4-amino-1H-pyrazolo[3,4-d]pyrimidin-3-yl)phenoxy)-4-methylphenyl)-4-(pyrrolidin-1-yl-methyl)-3-(trifluoromethyl)benzamide (6j). Prepared in the same manner as 6a. Yield: 34.9%, mp: 148–149 °C. ¹H NMR (300 MHz, DMSO-*d*₆) δ 13.58 (s, 1H), 10.49 (s, 1H), 8.24 (s, 2H), 8.20 (s, 1H), 7.94 (d, *J* = 8.0 Hz, 1H), 7.70 (d, *J* = 8.2 Hz, 2H), 7.61 (d, *J* = 9.1 Hz, 1H), 7.58 (s, 1H), 7.37 (d, *J* = 8.2 Hz, 1H), 7.12 (d, *J* = 8.2 Hz, 2H), 6.76 (s, 1H), 3.83 (s, 2H), 2.24 (s, 3H), 1.76 (s, 4H). HRMS (ESI+) *m/z* calculated for C₃₁H₂₈F₃N₇O₂ [M+H]⁺, 588.2329; found, 588.2334.

N-(3-(4-(4-amino-1H-pyrazolo[3,4-d]pyrimidin-3-yl)phenoxy)-4-methylphenyl)-4-(morpholino-methyl)-3-(trifluoromethyl)benzamide (6k). Prepared in the same manner as 6a. Yield: 37.9%, mp: 150–151 °C. ¹H

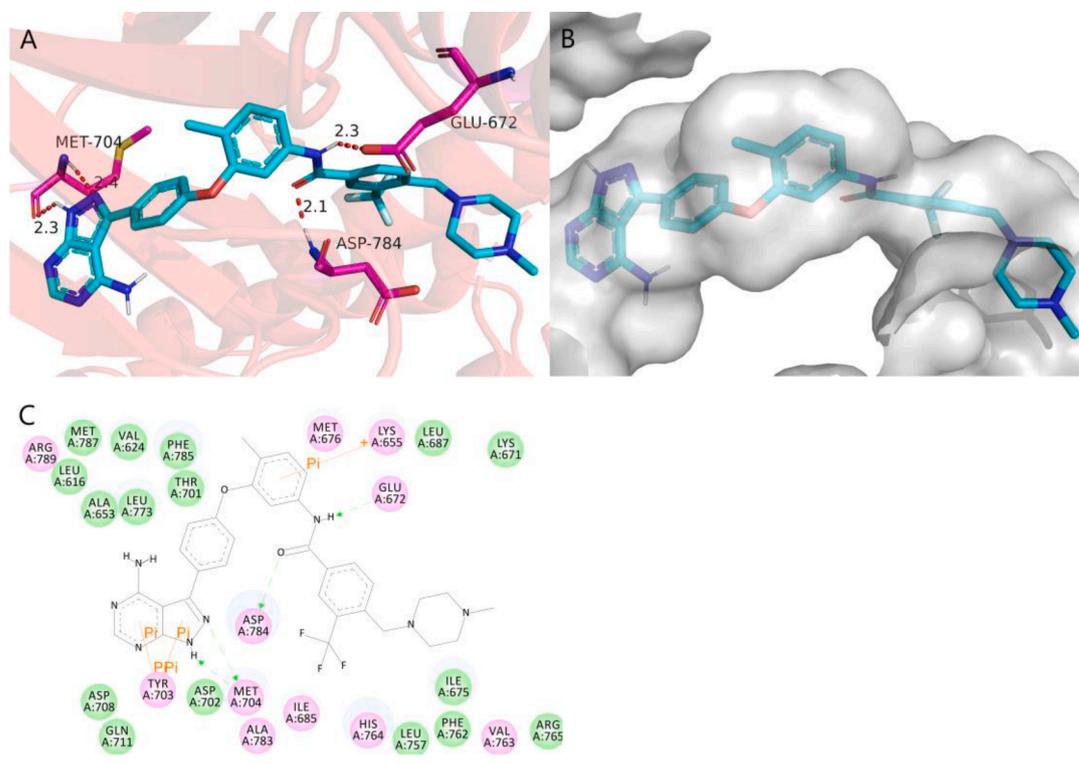


Fig. 3. Molecular docking of **6c** (blue) into the DDR1 (PDB ID: 4CKR). (A) Hydrogen-bond interactions between **6c** (in blue) and Met⁷⁰⁴, Glu⁶⁷², and Asp⁷⁸⁴, indicated by dash lines (B) Compound **6c** was fitted into the cavity of DDR1 kinase. (C) Compound **6c** interacted with residues in the DDR1 kinase domain. Residues in pink represented the electrostatic interactions with **6c**, and residues in green represented the Van der Waals interactions with **6c**.

NMR (300 MHz, DMSO-*d*₆) δ 13.56 (s, 1H), 10.47 (s, 1H), 8.21 (s, 3H), 7.93 (d, *J* = 8.4 Hz, 1H), 7.67 (d, *J* = 8.3 Hz, 2H), 7.57 (d, *J* = 9.9 Hz, 2H), 7.35 (d, *J* = 8.3 Hz, 1H), 7.09 (d, *J* = 8.4 Hz, 2H), 3.68 (s, 2H), 3.60 (s, 4H), 2.40 (s, 4H), 2.20 (s, 3H). HRMS (ESI⁺) *m/z* calculated for C₃₁H₂₈F₃N₇O₃ [M+H]⁺, 604.2278; found, 604.2282.

N-(3-(4-(4-amino-1H-pyrazolo[3,4-d]pyrimidin-3-yl)phenoxy)-4-methylphenyl)-4-(piperidin-1-yl-methyl)-3-(trifluoromethyl)benzamide (**6l**). Prepared in the same manner as **6a**. Yield: 35.6%, mp: 216–217 °C. ¹H NMR (300 MHz, DMSO-*d*₆) δ 13.59 (s, 1H), 10.50 (s, 1H), 8.28–8.17 (m, 3H), 7.96 (d, *J* = 8.2 Hz, 1H), 7.75–7.67 (m, 2H), 7.66–7.57 (m, 2H), 7.38 (d, *J* = 8.2 Hz, 1H), 7.17–7.09 (m, 2H), 6.78 (s, 1H), 3.66 (s, 2H), 2.39 (s, 4H), 2.24 (s, 3H), 1.63–1.38 (m, 6H). HRMS (ESI⁺) *m/z* calculated for C₃₂H₃₀F₃N₇O₂ [M+H]⁺, 602.2486; found, 602.2489.

N-(3-(4-(4-amino-1H-pyrazolo[3,4-d]pyrimidin-3-yl)phenoxy)-4-methylphenyl)-4-((4-methyl-piperidin-1-yl)methyl)-3-(trifluoromethyl)benzamide (**6m**). Prepared in the same manner as **6a**. Yield: 40.5%, mp: 242–243 °C. ¹H NMR (300 MHz, DMSO-*d*₆) δ 13.55 (s, 1H), 10.45 (s, 1H), 8.19 (t, 3H), 7.91 (d, *J* = 8.0 Hz, 1H), 7.67 (d, *J* = 8.2 Hz, 2H), 7.57 (d, *J* = 11.6 Hz, 2H), 7.34 (d, *J* = 8.2 Hz, 1H), 7.09 (d, *J* = 7.9 Hz, 2H), 3.63 (s, 2H), 2.73 (d, *J* = 10.0 Hz, 2H), 2.20 (s, 3H), 1.99 (t, *J* = 10.9 Hz, 2H), 1.57 (d, *J* = 12.5 Hz, 2H), 1.26–1.05 (m, 2H), 0.89 (d, *J* = 6.5 Hz, 3H). HRMS (ESI⁺) *m/z* calculated for C₃₃H₃₂F₃N₇O₂ [M+H]⁺, 616.2642; found, 616.2647.

N-(3-(4-(4-amino-1H-pyrazolo[3,4-d]pyrimidin-3-yl)phenoxy)-4-methylphenyl)-4-((4-ethyl-piperazin-1-yl)methyl)benzamide (**6n**). Prepared in the same manner as **6a**. Yield: 18.6%, mp: 276–277 °C. ¹H NMR (300 MHz, DMSO-*d*₆) δ 13.55 (s, 1H), 10.23 (s, 1H), 8.21 (s, 1H), 7.87 (d, *J* = 8.0 Hz, 2H), 7.69–7.64 (m, 2H), 7.60–7.55 (m, 2H), 7.42 (d, *J* = 8.0 Hz, 2H), 7.32 (d, *J* = 8.8 Hz, 1H), 7.11–7.06 (m, 2H), 3.51 (s, 2H), 2.46–2.23 (m, 10H), 2.19 (s, 3H), 0.97 (t, *J* = 7.2 Hz, 3H). HRMS (ESI+ESI⁺) *m/z* calculated for C₃₂H₃₄N₈O₂ [M+H]⁺, 563.2877; found, 563.2889.

3-(4-(4-amino-1H-pyrazolo[3,4-d]pyrimidin-3-yl)phenoxy)-*N*-(3-fluoro-4-((4-methylpiperazin-1-yl)methyl)phenyl)-4-methylbenzamide (**6o**). Prepared in the same manner as **6a**. Yield: 39.5%, mp: 148–150 °C. ¹H

NMR (300 MHz, DMSO-*d*₆) δ 13.60 (s, 1H), 10.34 (s, 1H), 8.24 (s, 1H), 7.80–7.67 (m, 4H), 7.63–7.53 (m, 3H), 7.36 (d, *J* = 8.8 Hz, 1H), 7.12 (d, *J* = 8.6 Hz, 2H), 3.59 (s, 2H), 2.42 (s, 4H), 2.36 (s, 4H), 2.23 (s, 3H), 2.17 (s, 3H). HRMS (ESI⁺) *m/z* calculated for C₃₁H₃₁FN₈O₂ [M+H]⁺, 567.2627; found, 567.2630.

3-(4-(4-amino-1H-pyrazolo[3,4-d]pyrimidin-3-yl)phenoxy)-*N*-(3-fluoro-4-(morpholinomethyl)-phenyl)-4-methylbenzamide (**6p**). Prepared in the same manner as **6a**. Yield: 44.0%, mp: 188–189 °C. ¹H NMR (300 MHz, DMSO-*d*₆) δ 13.56 (s, 1H), 10.31 (s, 1H), 8.21 (s, 1H), 7.78–7.70 (m, 2H), 7.70–7.63 (m, 2H), 7.61–7.50 (m, 3H), 7.33 (d, *J* = 8.7 Hz, 1H), 7.09 (d, *J* = 8.5 Hz, 2H), 3.56 (d, *J* = 5.5 Hz, 6H), 2.39 (t, *J* = 4.6 Hz, 4H), 2.20 (s, 3H). HRMS (ESI⁺) *m/z* calculated for C₃₀H₂₈FN₇O₃ [M+H]⁺, 554.2310; found, 554.2318.

3-(4-(4-amino-1H-pyrazolo[3,4-d]pyrimidin-3-yl)phenoxy)-*N*-(3-chloro-4-((4-methylpiperazin-1-yl)methyl)phenyl)-4-methylbenzamide (**6q**). Prepared in the same manner as **6a**. Yield: 44.0%, mp: 164–166 °C. ¹H NMR (300 MHz, DMSO-*d*₆) δ 13.56 (s, 1H), 10.34 (s, 1H), 8.21 (s, 1H), 7.98 (s, 1H), 7.86 (d, *J* = 8.0 Hz, 1H), 7.66 (d, *J* = 8.2 Hz, 2H), 7.63–7.54 (m, 3H), 7.33 (d, *J* = 8.2 Hz, 1H), 7.09 (d, *J* = 8.2 Hz, 2H), 3.60 (s, 2H), 2.44 (s, 4H), 2.33 (s, 4H), 2.20 (s, 3H), 2.16 (s, 3H). HRMS (ESI⁺) *m/z* calculated for C₃₁H₃₁ClN₈O₂ [M+H]⁺, 583.2331; found, 583.2340.

3-(4-(4-amino-1H-pyrazolo[3,4-d]pyrimidin-3-yl)phenoxy)-*N*-(3-chloro-4-(morpholinomethyl)-phenyl)-4-methylbenzamide (**6r**). Prepared in the same manner as **6a**. Yield: 61.0%, mp: 198–200 °C. ¹H NMR (300 MHz, DMSO-*d*₆) δ 13.58 (s, 1H), 10.37 (s, 1H), 8.24 (s, 1H), 8.02 (d, *J* = 1.7 Hz, 1H), 7.90 (dd, *J* = 8.0, 1.8 Hz, 1H), 7.73–7.57 (m, 5H), 7.36 (d, *J* = 8.3 Hz, 1H), 7.15–7.08 (m, 2H), 3.62 (dd, *J* = 8.0, 3.6 Hz, 6H), 2.45 (t, *J* = 4.6 Hz, 4H), 2.23 (s, 3H). ¹³C NMR (75 MHz, DMSO-*d*₆) δ 164.34, 158.54, 158.03, 156.52, 156.24, 153.89, 144.41, 139.37, 138.81, 135.47, 133.76, 131.87, 131.17, 130.49, 128.74, 128.32, 126.84, 125.01, 118.05, 116.81, 112.26, 97.38, 66.65, 59.24, 53.70, 15.88. IR (cm⁻¹): 3461.27, 3411.61, 3300.06, 3091.84, 2965.77, 2853.38, 2814.50, 1670.93, 1644.14, 1601.02, 1586.29, 1534.44, 1507.30,

1459.30, 1449.19, 1436.63, 1411.32, 1313.30, 1290.46, 1264.71, 1233.95, 1218.70, 1168.25, 1158.02, 1110.25, 1006.58, 990.06, 909.18, 864.18, 848.53, 814.36, 801.96, 740.84. HRMS (ESI+) m/z calculated for $C_{30}H_{28}ClN_7O_3$ $[M+H]^+$, 570.2015; found, 570.2022.

3-(4-(4-amino-1H-pyrazolo[3,4-d]pyrimidin-3-yl)phenoxy)-N-(3-bromo-4-((4-methylpiperazin-1-yl)methyl)phenyl)-4-methylbenzamide (6s). Prepared in the same manner as 6a. Yield: 44.0%, mp: 180~181 °C. 1H NMR (300 MHz, DMSO- d_6) δ 13.59 (s, 1H), 10.38 (s, 1H), 8.24 (s, 1H), 8.17 (s, 1H), 7.93 (d, J = 8.1 Hz, 1H), 7.69 (d, J = 8.2 Hz, 2H), 7.65–7.56 (m, 3H), 7.36 (d, J = 8.0 Hz, 1H), 7.11 (d, J = 8.2 Hz, 2H), 3.61 (s, 2H), 2.43 (s, 8H), 2.23 (s, 6H). HRMS (ESI+) m/z calculated for $C_{31}H_{31}BrN_8O_2$ $[M+H]^+$, 627.1826; found, 627.1823.

3-(4-(4-amino-1H-pyrazolo[3,4-d]pyrimidin-3-yl)phenoxy)-N-(3-bromo-4-(morpholinomethyl)-phenyl)-4-methylbenzamide (6t). Prepared in the same manner as 6a. Yield: 27.7%, mp: 189~190 °C. 1H NMR (300 MHz, DMSO- d_6) δ 13.59 (s, 1H), 10.38 (s, 1H), 8.24 (s, 1H), 8.19 (d, J = 1.7 Hz, 1H), 7.94 (dd, J = 8.0, 1.8 Hz, 1H), 7.75–7.56 (m, 5H), 7.36 (d, J = 8.1 Hz, 1H), 7.16–7.07 (m, 2H), 3.63 (d, J = 5.5 Hz, 6H), 2.46 (t, J = 4.6 Hz, 4H), 2.23 (s, 3H). HRMS (ESI+) m/z calculated for $C_{30}H_{28}BrN_7O_3$ $[M+H]^+$, 614.1510; found, 614.1513.

N-(3-(4-(4-amino-1H-pyrazolo[3,4-d]pyrimidin-3-yl)phenoxy)phenyl)-4-((4-methylpiperazin-1-yl)methyl)-3-(trifluoromethyl)benzamide (6u). Prepared in the same manner as 6a. Yield: 38.7%, mp: 146~148 °C. 1H NMR (300 MHz, DMSO- d_6) δ 13.55 (s, 1H), 10.51 (s, 1H), 8.23 (s, 3H), 7.92 (d, J = 8.0 Hz, 1H), 7.74–7.66 (m, 2H), 7.61 (dd, J = 8.5, 1.2 Hz, 2H), 7.42 (t, J = 8.3 Hz, 1H), 7.26–7.17 (m, 2H), 6.95–6.86 (m, 1H), 6.75 (s, 1H), 3.68 (s, 2H), 2.43 (s, 4H), 2.37 (s, 4H), 2.18 (s, 3H). HRMS (ESI+) m/z calculated for $C_{31}H_{29}F_3N_8O_2$ $[M+H]^+$, 603.2438; found, 603.2442.

4.3. In vitro kinase assay

The effects of compounds on the kinases DDR1 were assessed using a LanthaScreen Eu kinase activity assay technology (Invitrogen). Kinase reactions are performed in a 10 μ L volume in low-volume 384-well plates. The kinases in the reaction buffer consist of 50 mM HEPES pH 7.5, 0.01% Brij-35, 10 mM $MgCl_2$, and 1 mM EGTA; the concentration of the fluorescein-poly GAT substrate (Invitrogen) in the assay is 100 nM. Kinase reactions were initiated with the addition of 100 nM ATP in the presence of serials of dilutions of compounds. The reactions were allowed to proceed for 1 h at room temperature before a 10 μ L preparation of EDTA (20 mM) and Eu-labeled antibody (4 nM) in time-resolved TR-FRET dilution buffer are added. The final concentration of antibody in the assay well is 2 nM, and the final concentration of EDTA is 10 mM. The plate is allowed to incubate at room temperature for one more hour before the TR-FRET emission ratios of 665/340 nm were acquired on a PerkinElmer EnVision multilabel reader (Perkin-Elmer, Inc.). Data analysis and curve fitting were performed using GraphPad Prism 7 software.

4.4. Inhibition on cell proliferation by MTT assay

Adherent cells were plated in 96-well culture plates with cell density of 3000–4000 cells/well and treated with indicated compounds by adding 100 μ L of medium containing compounds of various concentrations on the second day. After 72 h of treatment, MTT was added to each well and incubated for additional 4–5 h, and the absorbance was measured on a microplate reader at 570 nm. Cell growth inhibition was evaluated as the ratio of the absorbance of the sample to that of the control.

4.5. Molecular docking assay

Molecular docking of the compound 6c into the three dimensional X-ray structure of DDR1 was carried out using the Discovery Studio (Version 4.0) as implemented through the graphical user interface

Discovery Studio CDOCKER protocol. The crystal structure of DDR1 (PDB ID: 4CKR) was obtained from the Protein Data Bank. All bound waters and ligands were eliminated from the protein and the polar hydrogens were added. The whole 4CKR was defined as a receptor and the binding site was selected based on ATP binding site of 4CKR. The three-dimensional structure of the compound 6c was constructed using ChemBio 3D Ultra 14.0 software and 6c was given CHARMM forcefield. The compound 6c was placed during the molecular docking procedure. Types of interactions of the docked protein with ligand were analyzed after the end of molecular docking.

Declaration of Competing Interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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