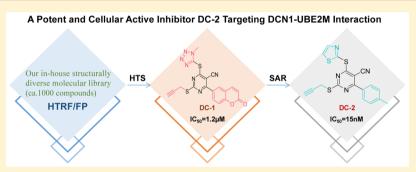
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Potent 5-Cyano-6-phenyl-pyrimidin-Based Derivatives Targeting DCN1-UBE2M Interaction

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Supporting Information



ABSTRACT: Neddylation of the Cullin-RING E3 ligases (CRLs) regulates the homeostasis of approximately 20% of cellular proteins. Defective in cullin neddylation 1 (DCN1), as a co-E3 ligase, interacts with UBE2M to enhance the activation of CRLs, and this interaction is emerging as a therapeutic target for human diseases. Here, we present a series of pyrimidin-based small molecular inhibitors targeting DCN1-UBE2M interaction. After finding a novel inhibitor DC-1 with IC₅₀ = 1.2 μ M, we performed a series of chemical optimizations, which finally led to the discovery of a potent thiazole containing 5-cyano-6phenylpyrimidin-based inhibitor DC-2 (IC₅₀ = 15 nM). Next, using protein and cellular thermal shift assays, coimmunoprecipitation, molecular docking, and site-specific mutation experiments, we further proved that DC-2 specifically inhibited the interaction of UBE2M and DCN1 at molecule and cellular levels, resulting in the decrease of cullin3 neddylation and accumulation of its substrate, NRF2. Our findings indicate that DC-2 may serve as a novel lead compound for specific derivatives targeting DCN1-UBE2M interaction.

INTRODUCTION

The ubiquitin-proteasome system (UPS) is integral to maintaining cellular protein homeostasis by regulating degradation of intracellular proteins. 1-7 The ubiquitylation pathway is executed by the coordinated efforts of the E1 (ubiquitin-activating enzyme), E2 (ubiquitin-conjugating enzyme), and E3 (ubiquitin ligase enzyme) proteins.⁵ Like ubiquitination,⁸ neddylation is a novel type of posttranslational modification, in which the ubiquitin-like molecule NEDD8 is added to its target proteins and thus regulates their functions. 9,10 Neddylation also consists of a tripartite enzymatic cascade, including the NEDD8-activating enzyme E1 (NAE, APPBP1 (NAE1), and UBA3 heterodimer), two NEDD8-conjugating enzymes E2s (UBE2F and UBE2M, also known as UBC12), and NEDD8-E3 ligases, 9,10 with cullins being the best-characterized substrates. 11-22 Up to now, neddylation has been recognized to be highly activated in various cancers, 20,27-33 which draws much attraction for oncologic drug discovery, especially on the finding of NEDD8 E1 inhibitor Pevonedistat (MLN4924). 10,23-26 MLN4924 covalently binds to NEDD8 E1, blocking all CRLs neddylation, thus causing accumulation of CRL substrates.^{7,10} Because MLN4924 possesses immense anticancer effects both in vitro and in vivo, 10,27-43 it has been approved for Phase II clinical trials for treatment of human acute myeloid leukemia, nonsmall cell lung cancer, and mesothelioma.²⁹ However, as a result of its broad ablation of neddylation, MLN4924 has a series of toxicities. Therefore,

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Figure 1. Representative examples of DCN1 inhibitors.

alternative targeting specific CRLs may have more potential and be safer for cancer treatment. $^{30-32}$

DCN1 (defective in cullin neddylation 1), also called DCUN1D1, DCNL1, or SCCRO, is a highly conserved gene and amplified along the 3q26.3 in most squamous cell carcinomas and some other human cancers. As a co-E3 ligase, it binds to the activation complex of Cullin-RBX1-UBE2M-NEDD8 to increase neddylation efficiency. Researchers have found that blocking UBE2M-DCN1 interaction can reduce cullin3 neddylation. Moreover, DCN1 overexpression has been reported to be associated with poor survival outcome, and DCN1 depression significantly reduces cancer cell growth and invasive capacity. 10,33-35,42,44-51 Therefore, DCN1 is considered as a promising and attractive anticancer target.

From the X-ray crystallographic structure of DCN1 with UBE2M, researchers have confirmed that during neddylation. the N-terminal acetylated UBE2M can dock into the hydrophobic pocket of DCN1, and this pocket could be used for designing DCN1 inhibitors.^{21,52} Up to now, four papers have reported the discovery of DCN1-UBE2M interaction inhibitors, ^{30-32,43} also known as DCN1 inhibitor. NAcM-COV, the first discovered inhibitor, showed highly specific effects on blocking the N-terminal acetylationdependent interaction of UBE2M with DCN1, resulting in the reduction of CUL1 and CUL3 neddylation in a squamous lung cancer cell line (HCC95).31,43 At the same time, Zhou et al, discovered two high-affinity inhibitors DI-591 and DI-404, which can selectively inhibit the neddylation of cullin 3 over other cullins, leading to the accumulation of cullin 3's substrate, NRF2 (Figure 1). 30,32 In this study, we screened our in-house structurally diverse molecular library (ca. 1000 compounds) and finally found a novel and potent pyrimidinebased DCN1-UBE2M interaction mediator DC-1, using both fluorescence polarization (FP) and homogeneous timeresolved fluorescence (HTRF)⁵³ assays. After extensive structure-activity relationship (SAR) efforts, a potent thiazole containing 5-cyano-6-phenylpyrimidin-based inhibitor DC-2 was discovered (Figure 2). DC-2 specifically targets DCN1-UBE2M interaction, leading to the inhibition of cullin3

$$SAR$$
 SAR
 SAR
 SAR
 SIR
 SIR

Figure 2. Discovery of potent thiazole contained pyrimidin-based inhibitor DC-2.

neddylation and the accumulation of NRF2 and NRF2' downstream proteins, HO-1 and NQO1. Our overall findings indicate that DC-2 may serve as a novel lead compound for specific derivatives targeting the DCN1-UBE2M interaction.

■ RESULTS AND DISCUSSION

Chemistry. The general synthesis route of the target pyrimidine-thiourea hybrids is depicted in Scheme 1. Benzaldehydes 1a-m, ethyl cyanoacetate 2, and thiourea 3 were prolonged heated in ethanol containing potassium carbonate to obtain 6-aryl-5-cyano-2-thiouracils 4a-m. Compounds 4a-m then reacted with the 3-bromoprop-1-yne, 3-bromoprop-1-ene, 1-bromopropane, and phosphorus oxychloride in dioxane to obtain the target derivatives 5a-o. S4 Compound 53 was prepared via click reaction of compound 5a with benzyl azide. After that, these highly activated intermediates (5a-o and 53) were reacted with appropriate mercapto heterocyclics and anilines to obtain compounds 6–52 and 54.

Development of FP- and HTRF-Based Competitive Binding Assays. To acquire stable and reliable binding affinities of our compounds in vitro, two methods (FP and HTRF) were used for screening DCN1–UBE2M interaction inhibitors. On the basis of the principle that the N-terminally acetylated UBE2M can interact with DCN1, ⁵⁵ UBE2M ^{NAc1-12} derivative fluorescently labeled tracer (FAM-782) and Histagged DCN1 recombinant protein were used to develope the fluorescence polarization (FP) assay. ³⁰ In addition, human GST-tagged DCN1 recombinant protein (Figure S1A) and AcUBE2M ¹⁻²¹-biotin peptides were applied to establish the HTRF screening system. The detailed schematic diagrams are shown in Figure 3A and B.

Biochemical Activity of the Candidate Compounds against DCN1–UBE2M Interaction and the Structure–Activity Relationship (SAR) Studies. All of the compounds synthesized in this study were examined for their inhibitory effects on DCN1–UBE2M interaction in vitro by both FP and HTRF assays. Compounds DI-591 and NAcM-COV were chosen as positive controls for FP³⁰ and HTRF assays, as respectively. All IC₅₀ values reported in this study were obtained from at least three independent experiments. The results are summarized in Tables 1–4.

To investigate the effect of the coumarin group on inhibitory activity, compounds 7-10 were synthesized initially, and their inhibition activities are shown in Table 1. As compared to the lead compound DC-1, compounds 7-9 with different heterocyclic substitution at R_1 caused a dramatic loss of activity. However, compound 10 with aromatic ring substitution performed with a more potent inhibitory effect. These findings indicate that the R_1 site plays a vital role in influencing their activities and the aromatic ring substitutions at R_1 may be

Scheme 1. Synthesis of the Target Pyrimidine-Based Derivatives

"Reagents and conditions: (a) absolute ethanol, absolute K_2CO_3 , reflux, 10 h, 70–90%; (b) (i) 3-bromoprop-1-yne, 3-bromoprop-1-ene, 1-bromopropane, dioxane, reflux; (ii) phosphorous oxychloride, reflux, 1 h, 50–80% (two steps); (c) benzyl azide, $CuSO_4$ - SH_2O , sodium ascorbate, $THF-H_2O$ (1:1), rt, 80%; (d) appropriate mercapto heterocyclic and aniline, absolute ethanol, reflux, 6 h, 70–95%.

A) FP assay for DCN1 binding to UBE2MNAc1-12 derivative



B) HTRF assay for DCN1 binding to UBE2MNAc1-21



Figure 3. Detailed schematic diagrams for the principle of screening DCN1–UBE2M interaction inhibitors. (A) FP assay for DCN1 binding to UBE2 $M^{NAc1-12}$ derivative. The FP value will be decreased after adding the compound, which can inhibit the DCN1–UBE2M interaction. (B) HTRF assay for DCN1 binding to UBE2 $M^{NAc1-21}$. Inhibitors will receive a low signal.

more important contributors in determining activity, as compared to those with coumarin group substitution.

On the basis of the above findings, further modifications were next focused on the substitution pattern and electronic effect on the aromatic ring of compound 10. As shown in Table 2, no matter an electron-withdrawing or -donating group substitution at R₁, the acquired compounds 11-17 display increased inhibitory activity as compared to no substitution compound 18. Compounds 11–14 with electron-withdrawing groups at R1 have a more potent inhibitory effect as compared to compounds 15-17 with electron-donating groups (except 10). The substitution of multiple functional groups on aromatic rings is beneficial to the improvement of activity (11 vs 12, 18 vs 17). Particularly, compound 14 with p-Br substitution effectively blocked the DCN1-UBE2M interaction with an IC₅₀ value of 65 nM, more potent than 12 with p-Cl substitution. These findings suggest that the number, type, and position of substituent and electronic effect on the aromatic ring have significant effects on the activity.

Next, through further structure—activity relationship (SAR) studies, we found that the heteroatoms and different heterocyclic substitutions at R₁ were also important for the inhibitory activity (Table 3). Replacing the sulfur atom at R₂ of compound 10 with nitrogen atom (19) led to a complete loss of activity. A similar trend was observed by changing the tetrazole-thione to other amino analogues (20-25). In contrast, sulfhydryl heterocyclic substitutions attached to the 4-position of the pyrimidine skeleton were well tolerated. As compared to 10, some of these compounds showed comparably or marginally increased potency, of which 34 with thiazolethiol group exhibited the most potent activity as compared to other heterocyclic substitutions (26-33). The extension of the side chain on nitrogen atom (26) has no significant effect on activity, as compared to the corresponding compound 10. In addition, we found that the number of nitrogen atoms on the mercapto heterocycles was important for the inhibitory activity: the tetrazole derivative 10 was more potent than the pyrazole derivative 27, and the triazole (28), pyridine (29) derivative without any detectable activity.

Table 1. Optimizing the Coumarin Group in the Structure of Compound 6 (DC-1) (Compounds 6-10)

$$N-N$$
 $N-N$
 $N-N$

C 1	R_1	DCN1-UBE2M	DCN1-UBE2M	
Compd.		$IC_{50} (\mu M)^a$ FP	$IC_{50} (\mu M)^a$ HTRF	
6	7,75	1.25 ± 0.08	0.82 ± 0.01	
(DC-1)	~~~~	1.20		
7	- \\ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \ \	> 20	>10	
8) NH	> 20	>10	
9		> 20	>10	
10	je do o	0.36 ± 0.01	0.25 ± 0.01	
DI-591		$20.82 \pm 1.24 \text{ nM}$	ND	
NAcM-COV		ND	$60.28 \pm 1.90 nM$	

 $^{{}^{}a}\text{IC}_{50}$ values were obtained from three independent repeats and represented as mean \pm SD. ND means the value was not determined.

Particularly, replacement of the nitrogen (N) atom in 28 with the sulfur (S) atom yielded 30, which significantly increased the inhibitory activity with an IC_{50} value of 195 nM. A similar trend was also observed in compounds 31–34, which suggested that the sulfur atom plays an important role in inhibitory activity. In addition, the introduction of hydrophilic functional groups (32, 33) reduced the inhibitory activity to some extent. Interestingly, as compared to 34, it was found that derivatizations of the olefinic bond on the thiazolethiol group (35–37) exhibited disappearance of activity, which indicated that the increase of steric hindrance at R_2 was detrimental to the activity.

To further investigate the importance of the two sites at R₁ and R₂ of 34, we simultaneously changed them and synthesized compounds 38-50. As compared to 34, some of these compounds showed comparable potency against DCN1, of which 42 (also named as DC-2) showed the most potent inhibitory effect on the DCN1-UBE2M interaction with an IC₅₀ value of 26 nM. In addition, compound 39 with chlorine substitution in the para position at the phenyl ring in R₁ displayed more potent inhibitory activity as compared to the meta- and disubstituted compounds (38 and 41). A similar trend was also observed (46 vs 47, 48), which suggested that the position of substituent at the phenyl ring in R1 plays an important role in the inhibitory activity. Furthermore, replacement of the phenyl group at R1 with aromatic heterocycle yielded 43-45, which significantly decreased the inhibitory activity, as well as 46-49 versus 50, that further suggested that the position of substituent and electronic effect at R₁ were important for the activity. Moreover, the

importance of terminal alkyne moiety was also evaluated. Because of the change of the propargyl group to ethene or ethyl group, compounds **51** and **52** exhibited no detectable (>10 000 nM) DCN1–UBE2M inhibitory effect as compared to **39**, which may be related to the hydrophobic interactions by forming π – π stacking with DCN1–UBE2M. In addition, replacing the propargyl group by triazole via click chemistry caused a dramatic loss of activity (**54** vs **39**), which indicated that the increase of steric hindrance at R₃ was detrimental to the activity. These modifications and SAR studies reveal that the terminal alkyne group is also critical for their inhibitory activity, and the introduction of hydrophobic functional groups of appropriate size at R₃ may be beneficial to the improvement of activity.

In summary, starting from a potent pyrimidine-based DCN1–UBE2M interaction mediator DC-1 (compound 6) in our in-house structurally diverse molecular library (ca. 1000 compounds), DC-2 (compound 42) containing 5-cyano-6-phenyl-pyrimidin was finally obtained with MW of 380, and two IC₅₀ values of 26.89 \pm 0.24 (K_i = 20.83 \pm 0.18 nM) and 15.71 \pm 1.19 nM (K_i = 13.66 \pm 1.03 nM) from the FP and HTRF assays (Figure 4A and 4B), respectively, resulting in about 80-fold improved potency. Furthermore, DC-2 has aqueous solubility in phosphate buffer (PBS, pH 7.4) to some extent (Figure S2). Therefore, DC-2 was chosen for the following target activity evaluation experiments. The compound 35 (DC-2N), which had a structure similar to DC-2 but presented no obvious inhibitory effect on DCN1–UBE2M interaction (IC₅₀ > 10 μ M), was chosen as a negative control.

Table 2. Optimization of the Substitution Pattern and Electronic Effect on the Aromatic Ring of Compound 10 (Compounds 11–18)

Compd.	D	DCN1-UBE2M	DCN1-UBE2M
	R_1	$IC_{50} (nM)^{a}$ -FP	$IC_{50} (nM)^{a}$ -HTRF
11	CI CI	78.49 ± 1.89	101.97 ± 2.00
12	, ye	644.13 ± 8.38	290.65 ± 2.31
13	Set CI	698.23 ± 6.48	281.65 ± 2.45
14	Br	64.75 ± 1.83	80.98 ± 1.90
15	res la constant de la	1089.11 ± 2.99	368.76 ± 2.22
16	reference to the second	3149.34 ± 5.87	1015.10 ± 3.00
17	7,25	942.93 ± 6.93	338.83 ± 2.53
10	3500	360.23 ± 3.64	251.13 ± 2.25
18	35	>20000	>20000

 $^{{}^}a\mathrm{IC}_{50}$ values were obtained from three independent repeats and represented as mean \pm SD. ND means the value was not determined.

DC-2 Binds to DCN1 and Increases Its Stability in Vitro. Next, the protein thermal shift assay was conducted to assess whether DC-2 can directly bind to DCN1 and adjust its stability in vitro. S6,57 As shown in Figure 5, as compared to DMSO control and DC-2N, DC-2 obviously caused an about 7 °C increase of the DCN1 melting temperature, similar to the effects of the positive control NAcM-COV. These findings indicate that DC-2 can directly bind to DCN1 and thus raise its thermal stability in vitro.

Binding Affinities of DC-2 on DCN Proteins. Previous studies have reported that there are five DCN1 homologues (DCNs: 1-5) in humans, ^{33,50} which contain a highly conserved central and C-terminal PONY domain, but possess distinct Nterminal regions. 33,50,58,59 The DCN1-5 PONY domains can interact with cullins to stimulate neddylation with different efficiencies. 55,60 Therefore, we further measured the binding affinities of DC-2 on DCN1-5 through competitive binding assay. As shown in Table 5, DC-2 also has a high binding affinity on DCN2, whose PONY domain in human shares 82% identity to that of DCN1.43 However, it shows relatively weak binding effects ($K_i > 500 \text{ nM}$) on DCN3-5, as compared to that of DCN1. These data suggest that DC-2 has specificity and selectivity for DCN1 and 2, similar to the previously reported inhibitors (DI-591, NAcM-COV, and DI-404).30,32,43

Furthermore, because the N-terminal domain of UBE2M has interaction with both DCN1 and NEDD8 E1 subunit UBA3, 52,61 we intended to determine whether **DC-2** also has inhibitory effects on E1's activity and UBE2M neddylation. As shown in Figure 6A and B, in the cell-free NAE activity assay, even though **MLN4924** significantly inhibits UBE2M neddylation, compound **DC-2** exhibits no obvious effect on NEDD8~UBE2M at concentration up to $100~\mu$ M, indicating that compound **DC-2** has no effect on E1's activity through ATP initiation and UBE2M neddylation.

Molecular Docking of DC-2 and DCN1. To investigate the binding mode of compound DC-2 with human DCN1, molecular docking study adopting the software MOE (The Molecular Operating Environment) version 2015.10 was carried out. Because the X-ray structure (PBD ID: 3TDU⁵²) obtained from the RSCB Protein Data Bank is the crystal structure of DCN1 in a complex with the N-terminal-acetylated UBE2M peptides and cullin 1, and our inhibitors were found on the basis of the principle that inhibitors compete with the N-terminal-acetylated UBE2M peptides to interact with DCN1, 3TDU was thus selected as the receptor to acquire the binding mode of DC-2 with DCN1. As shown in Figure 7A and B, compound DC-2 can be successfully docked into the active site of DCN1. There are 12 of 20 top-score docking conformations, which adopt a similar docking pose.

Table 3. Optimization of the Heteroatoms and Different Heterocyclic Substitutions of Compound 10 (Compounds 19-37)

$$R_2$$
 CN
 S
 N
 O
 O

		`	
Compd.	R_2	DCN1-UBE2M	DCN1-UBE2M
		IC ₅₀ (nM) ^a -FP	IC ₅₀ (nM) ^a -HTRF
19	N-N N'\ N N 252	>10000	>10000
20	HN N-{}-	>10000	>5000
21	S N-\{\xi	>10000	>5000
22	Ο N-ξ-	>10000	>10000
23	N-{-	>10000	>5000
24	N 1/2	>10000	>10000
25	F ₃ C N ² ²	>10000	>10000
26	N-N N S ² z ₂	591.11 ± 2.75	199.07 ± 2.29
27	N S 72	1217.90 ± 1.32	>5000
28	HN-N	>10000	>2500
29	N S	>10000	>2500
30	S S S S S S S S S S S S S S S S S S S	195.66 ± 2.29	65.98 ± 1.81
31	N-N Sixi	152.82 ± 2.18	133.60 ± 2.12
32	H ₂ N S S	820.78 ± 2.91	106.73 ± 2.02
33	H ₂ N S _z zz	413.55 ± 2.04	141.22 ± 2.15
34	S. S.	141.22 ± 2.10	77.04 ± 1.88
35 (DC-2N)	N S	>10000	>10000
36	N S	>10000	>10000
37	S S	>10000	>5000
10	N-N N'N S ³ 25	360 ± 3.64	251.13 ± 2.25

 $[^]a$ IC₅₀ values were obtained from three independent repeats and represented as mean \pm SD. ND means the value was not determined.

Table 4. Optimization of the Aromatic and Heterocyclic Substitutions at R₁, R₂, and R₃ of Compound 34 (Compounds 38-54)

$$R_{3}$$

Compd.	R_1	R_2	R ₃	DCN1-UBE2M	DCN1-UBE2M
Compa.	Kl	Κ2	Ν3	IC_{50} $(nM)^a$ -FP	$IC_{50} (nM)^{a}$ -HTRF
34	3400	S S	Soci.	141.22 ± 2.10	77.04 ± 1.88
38	; ^z ·CI	S S	Zr.	361.69 ± 2.93	140.38 ± 2.14
39	, zz CI	S sign	- Seri	36.59 ± 0.18	96.05 ± 1.98
40) St. Br	S sin	- Secti	551.87 ± 1.71	189.41 ± 1.95
41	, of CI	S Siri	- svr	62.99 ± 1.79	112.26 ± 2.05
42 (DC-2)	j. j. j.	S sin	- svr	26.89 ± 0.24	15.71 ± 1.19
43	34	S Sign	- Seri	789.09 ± 2.89	202.05 ± 2.00
44	rest N	S Sign	No.	1482.32 ± 1.39	1064.73±1.07
45	- '\\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\ \\	S sin	20th	>20000	>10000
46	, cf. CI	-N S 3 2 2	- sec	63.51 ± 1.80	122.65 ± 2.08
47	ř.ř. CI	-N N S 3 5	Zer.	106.21 ± 2.02	379.91 ± 2.58
48	ÇI CI	N-N N S 32	Sec.	109.04 ± 2.03	119.80 ± 2.07
49	ř	-N S-24	- sec	390.85 ± 2.56	172.29 ± 2.23
50	75	-N N S 3/2	200 Servi	665.64 ± 2.74	814.60 ± 2.91
51	7.F. CI	S Sign	Sold,	> 10000	> 10000
52	^z ^z CI	S Sign	~~~ strti	> 10000	> 10000
54	P. CI	S S	N,N,	> 10000	> 10000

 $[^]a$ IC $_{50}$ values were obtained from three independent repeats and represented as mean \pm SD. ND means the value was not determined.

Among them, the conformation with the highest score is illustrated in Figure 7A. The nitrogen atom of cyan group of DC-2 forms a hydrogen bond with the hydroxyl group of the side chain of Tyr181 with a distance of 2.09 Å. The sulfur atom connecting the thiazole ring of DC-2 forms a nontraditional hydrogen bond with the carbonyl group of the backbone of Pro97 with a distance of 3.55 Å. Besides, the methylbenzene group of DC-2 forms arene-H interaction with Phe164. The binding pocket surface of DCN1 and compound DC-2 is

shown in Figure 7B. The methylbenzene group of DC-2 is buried into the inside hydrophobic pocket surrounded by Phe117, Phe89, Ile86, Ala106, Val102, and Leu103. The propargyl group of DC-2 is located in the hydrophobic regions, which are formed by Gln87, Ile83, Ile86, and Cys115. In addition, the thiazole ring of DC-2 forms hydrophobic interactions with Tyr181, Leu184, and Met177. All of these computer-based predicted interactions indicate that compound

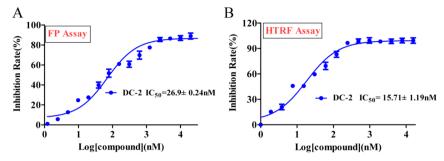


Figure 4. Compound DC-2 (compound 42) combines with DCN1 and thus inhibits UBE2M-DCN1 interaction in vitro. Compound DC-2 was added into the screening systems, and its inhibition rates (%) at different concentration were determined by both FP (A) and HTRF (B) assays, respectively. Data are presented as means \pm SD. Three individual experiments were performed for each group.

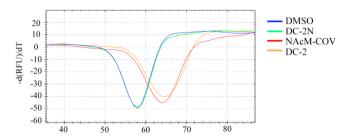


Figure 5. Compound DC-2 increases the stability of DCN1 in vitro. Normalized thermal shift assay response for recombinant human DCN1 in the presence of 0.5 mM DC-2 (orange) and DC-2N (green). NAcM-COV (red) was used as a positive control. DMSO (blue) was used as a blank control. Three individual experiments were performed for each group.

DC-2 could be well and specifically docked into the binding pocket of DCN1.

Site-Specific Mutations in DCN1 Binding Pockets with DC-2. Previous studies have reported that the Pro97, Phe164, and Tyr181 residues of human DCN1 are involved in the interaction of DCN1 and UBE2M, especially Tyr181, which can clamp the UBE2M's N-acetyl-Met1 and Ile2, pressing UBE2M's N-acetyl-Met into DCN1's hydrophobic pocket. 52,55 In addition, the reported DCN1-UBE2M interaction inhibitors all have interactions with these three residues, either by forming hydrogen bonds or by fitting into the hydrophobic pocket. 30,32,43 Therefore, we hypothesize that these three residues may play an important role in DCN1-UBE2M interaction. Furthermore, our above docking results also have confirmed that DC-2 has relatively strong interactions with Pro97, Phe164, and Tyr181 residues of DCN1 (Figure 7A and B). Therefore, we speculate that the inhibitory effects of compound DC-2 on DCN1-UBE2M interaction may be also related to these three amino acid residues. To prove our hypothesis, the three residues of DCN1 (Pro97, Phe164, and Tyr181) were chosen to do the sitespecific mutations to Thr97 (P97T), Ser164 (F164S), and Ile181 (Y181I), whose physicochemical properties are totally different from their original amino acids, respectively (Figure S1C). In our in vitro FP and HTRF assays, we found the complete loss of mp values and 665/615 ratios in DCN1

F164S and Y181I mutations, indicating that there is no interaction between the DCN1 mutations and AcUBE2M^{1–21}-biotin peptides. However, these parameters in DCN1 P97T mutation remained almost the same as those in wild-type DCN1, indicating that the P97T mutation did not affect its binding affinity with AcUBE2M^{1–21}-biotin (Figure 7C and D). To confirm this discovery, we further did the label-free BioLayer Interferometry (BLI) assay. Consistent with our above findings, DCN1 F164S and Y181I mutations lost the binding affinity with AcUBE2M^{1–21}-biotin peptide rather than DCN1 P97T mutation (Figure S3). All of these findings suggest that both Phe164 and Tyr181 residues are crucial for DCN1–UBE2M interaction, and the inhibition effect of DC-2 may be related to its binding interaction with these two residues.

DCN1 Was Highly Activated in Several Human Cancer Cell Lines. To choose suitable cancer cell lines for the further experiments, the DCN1 protein levels in several human cancer cell lines, including lung cancer cell lines (PC9, A549, and H1975), esophageal cancer cell lines (EC109, EC9706, KYSE70, KYSE140, and TE-1), liver cancer cell lines (SMMC-7721, BEL-7402, and ZIP177), prostatic cancer cell line (PC3), and breast cancer cell line (MCF7), were measured. We found that DCN1 was highly expressed in these cancer cell lines, as compared to those in normal cell lines: GES-1, Het-1A, and L02, respectively (Figure 8A–C).

Interaction of DC-2 with DCN1 at the Cellular Level. To further determine whether DC-2 can target intracellular DCN1, cellular thermal shift assay (CETSA) was performed in one of the DCN1 overexpressed cancer cell line H1975. As shown in Figure 9A and D, DCN1 protein started to degrade at 53 °C in DMSO or 10 μ M DC-2N treated cells (Figure 9A,B,D, and E), while it was very stable from 43 to 61 °C in the 10 μ M DC-2 treated cells, similar to that treated by positive compound DI-591 (Figure 9A and D).³⁰ In addition, DC-2 increases DCN1 thermal ability at a concentration low to 1 μ M at 55 °C (Figure 9B and E). These findings indicate that DC-2 can engage the cellular DCN1 protein and increase its thermal ability. Furthermore, the DCN1 level, which was pulled down by UBE2M antibody, was obviously decreased in DC-2 treated cells, as compared to those in DMSO or DC-2N treated groups (Figure 9C and F). These findings further

Table 5. Binding Affinities of Compound DC-2 on the Indicated DCN1-5 Recombinant Proteins

	DCN1 K_i $(nM)^a$	DCN2 K_i $(nM)^a$	DCN3 K_i $(nM)^a$	DCN4 $K_i (nM)^a$	DCN5 $K_i (\mu M)^a$
DC-2	13.66 ± 1.03	66.19 ± 1.72	591.72 ± 5.17	807.01 ± 9.42	2.14 ± 0.32

 $^{{}^{}a}K_{i}$ values were obtained from three independent repeats and represented as mean \pm SD. ND means the value was not determined.

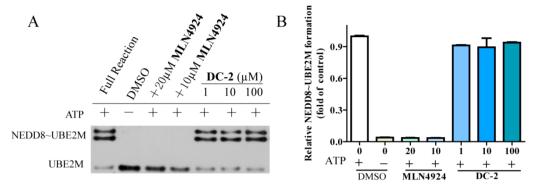


Figure 6. Compound DC-2 exhibits no obvious inhibitory effect on UBE2M neddylation level in vitro. (A) After adding 10 and 20 μ M MLN4924 or 1, 10, and 100 μ M DC-2 into the NEDD8, NEDD8 E1, and UBE2M complex, respectively, the NEDD8~UBE2M formation was initiated by ATP. (B) Densitometry shows relative protein expression level, which was analyzed by fold of DMSO with ATP group. Data are presented as means \pm SD. Three individual experiments were performed for each group.

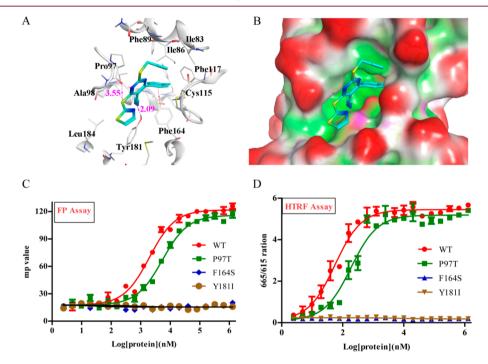


Figure 7. Predicted binding mode of compound DC-2 in DCN1 binding pocket (PDB ID: 3TDU). (A) Residues forming interactions in the docked complex. Compound DC-2 is shown as cyan stick; the residues associated with the compound are shown as white lines. Hydrogen bonds are shown as magenta dash lines, and the arene—H interaction is shown as a red dash line. The corresponding distances are given in angstroms. (B) The binding pocket surface of DCN1 and compound DC-2. Green area represents hydrophobic region, and red area represents exposed solvent region. The mp values (C) and 665/615 ratios (D) were determined after adding increased concentration of recombinant wild-type (WT) DCN1 or its site-specific mutations: P97T, F164S, and Y181I. Data are presented as means \pm SD. At least three individual experiments were performed for each group.

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suggest that compound DC-2 can bind to DCN1 and inhibit the association of UBE2M and DCN1 at the cellular level.

Selectively Blocking the Neddylation of Cullin3 by DC-2 in Lung Cancer Cells. Human DCN1 acts as a co-E3 ligase, which can interact with cullins and Rbx1 to stimulate the neddylation of cullins in vivo, especially cullin1—4. ^{21,40,52,55,60,62} To examine the effect of DC-2 on the neddylation of cullins at cellular level, two DCN1 highly expressed lung cancer cell lines H1975 and PC9 were treated with compound DC-2 for 24 h, using DC-2N or MLN4924 as a negative or positive control, respectively. We found that DC-2 treatment resulted in a significantly decreased cullin3 (CUL3) neddylation level, without obvious impact on the neddylation of other cullins (CUL1, CUL2, CUL4A, CUL4B,

and CUL5, Figure 10A) in both cell lines. In addition, because the inhibition of the cullins neddylation has been associated with the accumulation of its substrates, the expression levels of nuclear factor-erythroid 2 related factor 2(NRF2) (one substrate of cullin3), 63-65 as well as NRF2's downstream proteins (NADPH: quinone oxidoreductase-1, NQO1) and (heme oxygenase-1, HO-1), were then determined after DC-2 treatment. 66-68 We found that the expression levels of NRF2, HO-1, and NQO1 were obviously increased in both H1975 and PC9 cells, similar to the findings treated by MLN4924 (MLN, Figure 10C). Yet the expression levels of CylinE1, p21 (two substrates of cullin1), and CDT1 (a substrate of cullin4A) were not changed after DC-2 treatment (Figure 10C).

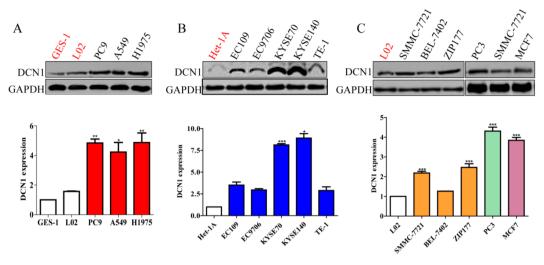


Figure 8. Expression levels of DCN1 in several human cancer cell lines. (A) Lung cancer cell lines: PC9, A549, and H1975. (B) Esophageal cancer cell lines: EC109, EC9706, KYSE70, KYSE140, and TE-1. (C) Liver cancer cell lines: SMMC-7721, BEL-7402, and ZIP177. Prostate cancer cell line: PC3. Breast cancer cell line: MCF7. Data are presented as means \pm SD. At least three individual experiments were performed for each group. *P < 0.05, **P < 0.01, ***P < 0.001 as compared to the normal control cell lines: GES-1, Het-1A, or L02, respectively.

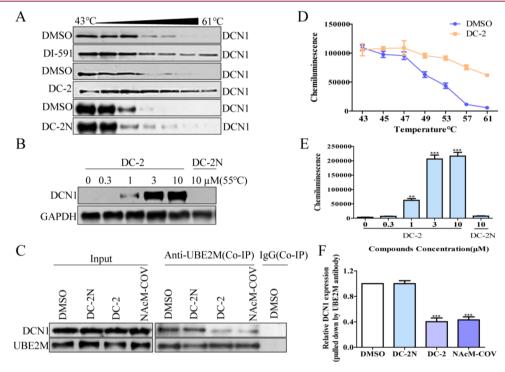


Figure 9. Compound DC-2 specifically binds to DCN1 and blocks the interaction of DCN1–UBE2M at a cellular level. (A and B) H1975 cells were treated with 10 μ M compound DC-2, DI-591, or DC-2N for 5 h, respectively. They then were collected and heated from 43 to 61 °C (A). H1975 cells were treated with 0, 0.3, 1, 3, 10 μ M DC-2 or 10 μ M DC-2N before being heated at 55 °C. Their DCN1 expression levels then were determined by Western Blot (B). (C) Compound DC-2, but not DC-2N, blocks the association of DCN1 and UBE2M at the cellular level. After H1975 cells were treated with compound DC-2N, DC-2, or NAcM-COV, the protein levels of UBE2M and DCN1, which were pulled down by UBE2M antibody, were determined by Western Blot (right panel). The basal levels of DCN1 and UBE2M in cell lysates were determined by Western Blot (left panel). (D–F) The bands intensities of proteins in (A)–(C), respectively. At least three individual experiments were performed for each group.

neddylation of cullin3 (Figure 10B, E, G, and H) as well as the accumulation of NRF2, NQO1, and HO-1 (Figure 10D, F, G, and H) exhibited in a dose (0.3, 1, 3, 10, and 20 μ M) and time-dependent manner (0.5, 1, 3, 12, 24, and 48 h), without significant effect on the neddylation of other cullins (CUL1–5) (Figure 10B) and their conresponding substrates: CylinE1, p21, and CDT1 (Figure 10D and G) in both H1975 and PC9

cells. These findings indicate the specifically inhibitory effects of DC-2 on the neddylation of cullin3 at the cellular level, without significant effects on other cullins neddylation.

Moreover, silencing DCN1 by three small interfering RNA (siRNA) in both PC9 and H1975 cells also decreased the neddylation of cullin3 and caused the accumulation of NRF2, NQO1, and HO-1 (Figure 11A–C). Furthermore, DC-2

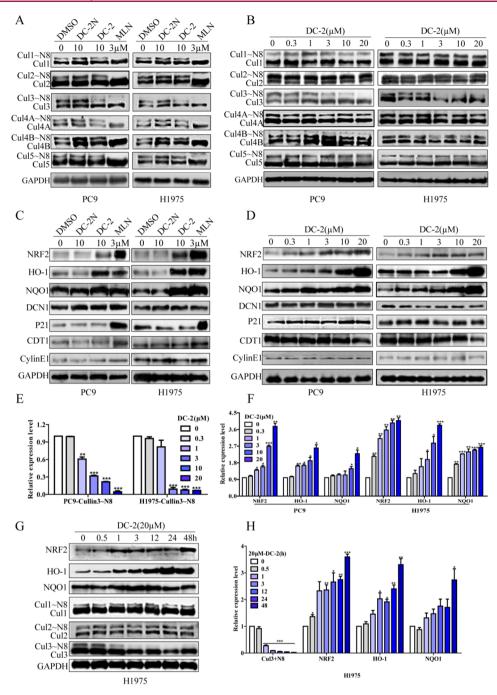


Figure 10. Compound **DC-2** inhibits intracellular cullin3's neddylation process and subsequently increases its substrate NRF2 as well as NRF2' downstream proteins: HO-1 and NQO1 with time (G and H) and dose (A–F)-dependent manner. Densitometry shows the relative protein expression level normalized by GAPDH. Data are presented as means \pm SD. Three individual experiments were performed for each group. *P < 0.05, **P < 0.01, ***P < 0.001 as compared to the DMSO controls.

treatment did not have an additive effect on them (Figure 11D–F). These results further confirm that the effect of DC-2 on the neddylation of cullin3 is through its specific binding to DCN1.

Effects of DC-2 on DCN1 Highly Expressed Cancer Cells. DCN1 is highly expressed in most cancers and has been discovered as an oncogene. ^{10,23,33-39,44-46,69} To determine whether DC-2 has the ability of inhibiting cancer cells' proliferation, eight DCN1 amplified cancer cell lines (KYSE70, PC9, SMMC-7721, PC3, MCF7, TE-1, KYSE140, and EC109) and two normal cell lines (GES-1 and L02) were chosen to be treated with DC-2 for 72 h. As shown in Figure 12A, DC-2

shows strong antiproliferation ability on these cancer cells rather than normal cells. Furthermore, compound DC-2 could decrease three types of lung cancer cells' viabilities (H1975, PC9, and A549) with time-dependent manner (Figure 12B), block their colony formations (Figure 12C and D), and induce their apoptosis dose dependently (Figure 12E and F). These results suggest that compound DC-2 has the ability of blocking DCN1 highly expressed cancer cells' proliferation. In addition, the cytotoxicities of DC-2N and the compounds with very similar structures of DC-2N (27, 29, 36, 37, and 45) or DC-2 (33, 39, 40, 46, and 47) on five DCN1 highly expressed cancer cells (KYSE70, PC9, SMMC-7721, PC3, and MCF7) were

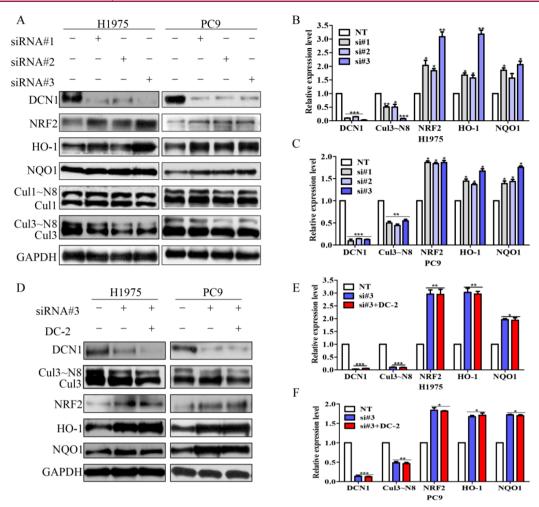


Figure 11. Protein levels of DCN1, NRF2, HO-1, NQO1, Cullin1, and Cullin3 were measured after H1975 and PC9 cells were treated with three DCN1 siRNA (A). Nontargeting siRNA (NT) treatment was used as control. (D) The protein levels of Cullin3, DCN1, NRF2, HO-1, and NQO1 were determined after treating the H1975 and PC9 cells by siRNA#3 with or without compound DC-2. (B, C, E, and F) Densitometry shows relative protein expression level normalized by GAPDH. Data are presented as means \pm SD. Three individual experiments were performed for each group. *P < 0.05, **P < 0.01, ***P < 0.001 as compared to the controls (NT).

determined. As shown in Table 6, compounds 33, 39, 40, 46, and 47, which can block the DCN1-UBE2M interaction at nanomole level in vitro, also have cytotoxicities on those tested cancer cells with IC₅₀ values ranging from 1 to 12 μ M. However, DC-2N, 27, 29, 36, 37, and 45, which have no inhibiting effect on DCN1-UBE2M interaction, exhibits less cytotoxic activity (>20 μ M), indicating that the cytotoxicities of these compounds may be related to their blocking effects on UBE2M-DCN1 interaction. However, the cytotoxic activities of DI-591 on the five DCN1 highly expressed cancer cell lines were also measured. The IC₅₀ values were all above 20 μ M at 72 h. On the basis of the current findings, whether DCN1 can be recognized as a valuable antitumor target remains to be further investigated. In addition, except for the inhibitory effects of DC-2 on DCN1-2, DC-2 also exhibits relatively weak binding affinities on DCN3-5 (Table 5), similar to the findings on NAcM-COV. 43 Therefore, the cytotoxicities of DC-2 may be also related to its inhibition on DCN3-5. However, the relative less cytotoxicity of DI-591 indicates its higher selectivity on DCN1-2 over DCN3-5.30

CONCLUSIONS

In this study, a novel series of compounds with a 5-cyano-6phenylpyrimidin scaffold as DCN1-UBE2M interaction modulators have been identified through structure-based optimization, which enriches the structure types of DCN1 inhibitors. Among the inhibitors, compound DC-2 exhibits the most potent inhibition effect on DCN1-UBE2M interaction at molecule and cellular levels. Molecular docking results show that DC-2 can well dock into the binding pocket of DCN1. Site-specific mutations further verified its blocking effects. Furthermore, unlike MLN4924, obliterating all cullins neddylation, DC-2 specifically diminishes the neddylation of cullin3, which leads to the accumulation of cullin3's substrate NRF2 and NRF2's downstream proteins: HO-1 and NQO1. Our findings indicate that the 5-cyano-6-phenylpyrimidinbased small molecules may serve as leading compounds specifically targeting DCN1-UBE2M interaction.

■ EXPERIMENTAL SECTION

General Methods for Chemistry. Chemicals and solvents were obtained from standard suppliers and used directly without further purification. Melting points were taken on an X-5 micromelting apparatus and were uncorrected. ¹H and ¹³C NMR spectra were

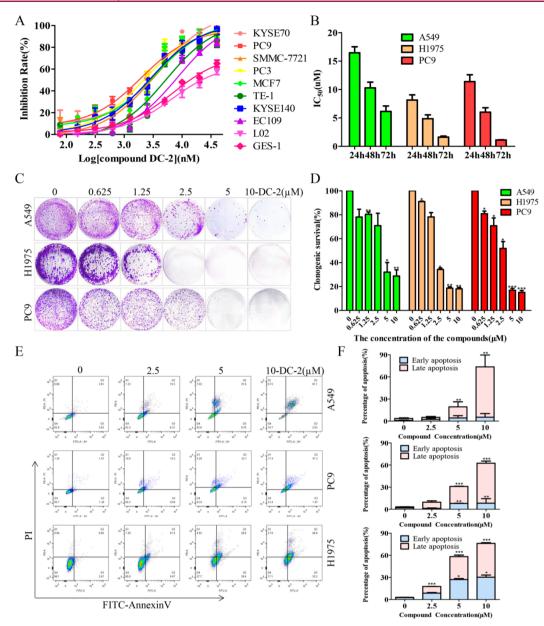


Figure 12. Compound DC-2 inhibits cell proliferation and colony formation. (A) The inhibition rates (%) of DC-2 on eight different cancer cell lines and two normal cell lines: L02 and GES-1. (B) The IC $_{50}$ values were determined after treating the three lung cancer cell lines (PC9, H1975, and A549) with DC-2 for 24, 48, and 72 h. (C) After the treatment of compound DC-2 at 0, 0.625, 1.25, 2.5, 5, and 10 μ M for 7 days, cells were stained by crystal violet and then imaged by microscopy. (D) Subsequently, they were dissolved and then measured by a BioTek microplate reader. (E and F) The apoptosis was determined by flow cytometry after PC9, H1975, and A549 cells were treated with compound DC-2 for 48 h. Data are presented as means \pm SD. Three individual experiments were performed for each group. *P < 0.05, **P < 0.01, ***P < 0.001 as compared to the controls.

respectively determined with a 400 and 100 MHz spectrometer. High-resolution mass spectra (HRMS) were obtained with a Water Q-TOF electrospray mass spectrometer (Water, Milford, MA). Final products were of >95% purity as analyzed by HPLC analysis (Phenomenex column, C-18, 5.0 μ m, 4.6 mm \times 150 mm) on a Dionex UltiMate 3000 UHPLC instrument from ThermoFisher. Besides, PAINS screening of the synthesized compounds was carried out by employing the online program, 70 and all of the tested compounds passed the filter.

General Procedure for the Synthesis of Compounds 6–52 and 54. To a well-stirred solution of the appropriate mercapto heterocyclics and anilines (5 mmol) in absolute ethanol (10 mL), an equimolar amount of a solution of compounds 5a–o or 53 (5 mmol) in absolute ethanol (10 mL) was added. The reaction mixture was stirred and heated under reflux for 5 h. Upon completion, the

precipitated product was filtered off and washed with ethanol to afford the crude product. The crude product was recrystallized from ethanol to yield the pure products 6-52 and 54. The detailed information on the synthesis and characterization of compounds 5a-o and 53 was reported in published articles. $^{54,71-73}$

4-((1-Methyl-1H-tetrazol-5-yl)thio)-6-(2-oxo-2H-chromen-6-yl)-2-(prop-2-yn-1-ylthio)pyrimidine-5-carbonitrile (6). Yield 88.5%. White solid. Mp: 184–185 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 8.36 (d, J=1.7 Hz, 1H, Ar–H), 8.22 (dd, J=6.2, 4.0 Hz, 2H, Ar–H), 7.66 (d, J=14.7 Hz, 1H, Ar–H), 6.62 (d, J=9.6 Hz, 1H, Ar–H), 4.14 (s, 3H, –CH₃), 3.57 (d, 2H, –CH₂–), 3.17 (s, 1H, J=2.5 Hz, \equiv C–H). 13 C NMR (100 MHz, DMSO- d_6 , ppm): δ 172.31, 169.73, 165.20, 159.34, 155.73, 145.48, 143.73, 132.19, 130.26, 129.64, 119.06, 117.39, 117.10, 114.61, 98.65, 78.79, 73.56, 34.79,

Table 6. Cytotoxic Activities (IC₅₀'s) of DC-2 and DC-2N Derivatives against Five DCN1 Highly Expressed Cancer Cell Lines^a

			$IC_{50} (\mu M)^b$			
compd	KYSE70	PC9	SMMC-7721	PC3	MCF7	DCN1-UBE2M IC ₅₀ (nM)-FP
27	>20	>20	>20	>20	>20	>10000
29	>10	>20	>20	>10	>10	>10000
DC-2N	>20	>20	>20	>20	>20	>10000
36	>20	>20	>20	>20	>20	>10000
37	>20	>20	>20	>20	>20	>10000
45	>20	>20	>20	>20	>20	>20000
33	8.25 ± 0.91	7.96 ± 0.90	4.66 ± 0.66	10.28 ± 1.01	9.82 ± 0.99	195.66 ± 2.29
39	5.35 ± 0.72	4.13 ± 0.61	6.04 ± 0.78	3.62 ± 0.55	3.23 ± 0.51	36.59 ± 0.18
40	10.88 ± 1.03	12.11 ± 1.08	3.99 ± 0.60	8.53 ± 0.93	11.63 ± 1.06	551.87 ± 1.71
DC-2	4.30 ± 0.63	1.11 ± 0.04	2.45 ± 0.39	1.62 ± 0.32	2.20 ± 0.34	26.89 ± 0.24
46	6.56 ± 0.81	5.30 ± 0.72	7.34 ± 0.86	7.17 ± 0.85	6.18 ± 0.79	63.51 ± 1.80
4 7	5.32 ± 0.72	6.48 ± 0.81	8.64 ± 0.93	3.54 ± 0.55	4.12 ± 0.61	106.21 ± 2.02
NAcM-COV	17.89 ± 1.25	5.59 ± 0.74	16.24 ± 1.21	25.27 ± 1.40	10.17 ± 1.00	ND

^aCells were treated with different concentrations of the indicated compounds for 72 h. Cell viability was measured by MTT assay as described in the Experimental Section. ${}^{b}IC_{50}$ values were indicated as the mean \pm SD of three independent experiments. ND means the value was not determined.

19.12. HR–MS (ESI) calcd for $C_{19}H_{11}N_7NaO_2S_2$: [M + Na]⁺ m/z, 456.0313; found, 456.0315.

4-((1-Methyl-1H-tetrazol-5-yl)thio)-2-(prop-2-yn-1-ylthio)-6-(thiophen-3-yl)pyrimidine-5-carbonitrile (7). Yield 83.9%. White solid. Mp: 183–184 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 8.32 (s, 1H, Ar–H), 8.13 (d, J = 4.1 Hz, 1H, Ar–H), 7.39 (s, 1H, Ar–H), 4.11 (s, 3H, –CH₃), 3.59 (d, 2H, –CH₂–), 3.15 (t, 1H, ≡C–H). ¹³C NMR (100 MHz, DMSO- d_6 , ppm): δ 171.87, 169.80, 158.29, 145.45, 138.13, 135.80, 132.60, 129.72, 114.96, 94.27, 78.77, 73.52, 34.75, 19.06. HR–MS (ESI) calcd for C₁₄H₉N₇NaS₃: [M + Na]⁺ m/z, 393.9979; found, 393.9982.

4-(1*H*-Indol-3-yl)-6-((1-methyl-1*H*-tetrazol-5-yl)thio)-2-(prop-2-yn-1-ylthio)pyrimidine-5-carbonitrile (8). Yield 86.6%. White solid. Mp: 227–227 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 12.37 (s, 1H, NH, D₂O exchangeable), 8.63 (d, J = 3.0 Hz, 1H, -CH=), 8.41 (d, J = 7.8 Hz, 1H, Ar-H), 7.57 (d, J = 7.9 Hz, 1H, Ar-H), 7.29 (t, J = 7.3 Hz, 1H, Ar-H), 7.24 (t, J = 7.5 Hz, 1H, Ar-H), 4.13 (s, 3H, -CH₃), 3.59 (d, J = 2.3 Hz, 2H, -CH₂-), 3.17 (t, J = 2.4 Hz, 1H, \equiv C-H). ¹³C NMR (100 MHz, DMSO- d_6 , ppm): δ 171.68, 168.60, 161.76, 136.52, 132.66, 125.44, 123.43, 122.29, 122.06, 116.14, 112.60, 110.30, 93.65, 79.15, 73.57, 34.75, 18.78. HR-MS (ESI) calcd for C₁₈H₁₂N₈NaS₂: $[M + Na]^+$ m/z, 427.0524; found, 427.0525.

4-((1-Methyl-1H-tetrazol-5-yl)thio)-2-(prop-2-yn-1-ylthio)-6-(1H-pyrrol-2-yl)pyrimidine-5-carbonitrile (9). Yield 51.6%. White solid. Mp: 189–190 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 12.08 (s, 1H, NH, D₂O exchangeable), 7.49 (s, 1H, -CH=), 7.29 (s, 1H, -CH=), 6.53–6.32 (m, 1H, -CH=), 4.10 (s, 3H, -CH₃), 3.69 (d, J = 2.6 Hz, 2H, -CH₂-), 3.15 (t, J = 2.5 Hz, 1H, \equiv C-H). 13 C NMR (100 MHz, DMSO- d_6 , ppm): δ 171.53, 168.77, 155.43, 145.62, 127.32, 125.45, 117.04, 115.44, 112.13, 91.76, 79.03, 73.67, 34.72, 18.86. HR–MS (ESI) calcd for C₁₄H₁₀N₈NaS₂: [M + Na]⁺ m/z, 418.0521; found, 418.0521.

4-((1-Methyl-1H-tetrazol-5-yl)thio)-2-(prop-2-yn-1-ylthio)-6-(3,4,5-trimethoxyphenyl)pyrimidine-5-carbonitrile (10). Yield 82.1%. White solid. Mp: 256–256 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 7.40 (s, 2H, Ar–H), 4.12 (s, 3H, –CH₃), 3.87 (s, 6H, –CH₃), 3.80 (s, 3H, –CH₃), 3.65 (t, J=12.7 Hz, 2H, –CH₂–), 3.20 (t, J=2.3 Hz, 1H, \equiv C–H). ¹³C NMR (100 MHz, DMSO- d_6 , ppm): δ 172.55, 170.08, 166.35, 153.39, 146.06, 141.54, 129.54, 115.48, 107.34, 98.83, 79.69, 73.84, 60.78, 56.69, 35.24, 19.77. HR–MS (ESI) calcd for C₁₉H₁₇N₇NaO₃S₂: [M + Na]⁺ m/z, 478.0732; found, 478.0733.

4-(4-Chlorophenyl)-6-((1-methyl-1H-tetrazol-5-yl)thio)-2-(prop2-yn-1-ylthio)pyrimidine-5-carbonitrile (11). Yield 87.7%. White solid. Mp: 163–164 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 8.02 (d, J = 8.4 Hz, 2H, Ar–H), 7.74 (t, J = 19.1 Hz, 2H, Ar–H), 4.13 (s, 3H, –CH₃), 3.59 (s, 2H, –CH₂–), 3.15 (s, J = 2.5 Hz, 1H, \equiv C–

H). 13 C NMR (100 MHz, DMSO- d_6 , ppm): δ 172.30, 169.69, 165.47, 145.47, 137.36, 132.92, 130.85, 129.06, 114.55, 98.72, 78.78, 73.51, 34.78, 19.10. HR–MS (ESI) calcd for $C_{16}H_{10}ClN_7S_2$: [M + Na]⁺ m/z, 422.0025; found, 422.0025.

4-(3-Chlorophenyl)-6-((1-methyl-1H-tetrazol-5-yl)thio)-2-(prop2-yn-1-ylthio)pyrimidine-5-carbonitrile (12). Yield 89.7%. White solid. Mp: 163–165 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 8.00 (t, J=1.8 Hz, 1H, Ar–H), 7.94 (d, J=7.8 Hz, 1H, Ar–H), 7.78–7.73 (m, 1H, Ar–H), 7.67 (t, J=7.9 Hz, 1H, Ar–H), 4.13 (s, 3H, –CH₃), 3.60 (d, J=7.5 Hz, 2H, –CH₂–), 3.16 (t, J=2.5 Hz, 1H, \equiv C–H). ¹³C NMR (100 MHz, DMSO- d_6 , ppm): δ 172.39, 169.63, 165.23, 145.46, 136.11, 133.55, 131.96, 130.87, 128.65, 127.65, 114.43, 99.11, 78.81, 73.46, 34.77, 19.12. HR–MS (ESI) calcd for C₁₆H₁₀ClN₇NaS₂: [M + Na]⁺ m/z, 422.0025; found, 422.0025.

4-(4-Bromophenyl)-6-((1-methyl-1H-tetrazol-5-yl)thio)-2-(prop2-yn-1-ylthio)pyrimidine-5-carbonitrile (13). Yield 78.1%. Yellow solid. Mp: 238–239 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 8.23 (d, J = 3.3 Hz, 1H, -CH \Longrightarrow), 8.13 (d, J = 3.3 Hz, 1H, -CH \Longrightarrow), 7.97–7.91 (m, 2H, Ar \Longrightarrow H, 7.88–7.82 (m, 2H, Ar \Longrightarrow H, 3.79 (d, J = 2.6 Hz, 2H, -CH \Longrightarrow DMSO- d_6 , ppm): δ 172.03, 171.48, 165.43, 150.29, 144.31, 133.50, 131.91, 131.04, 127.91, 126.18, 114.45, 98.20, 79.06, 73.86, 19.29. HR \Longrightarrow MS (ESI) calcd for C $_{16}$ H $_{10}$ BrN $_7$ NaS $_2$: [M + Na] $^+$ m/z, 466.9070; found, 466.9073.

4-((1-Methyl-1H-tetrazol-5-yl)thio)-2-(prop-2-yn-1-ylthio)-6-(p-tolyl)pyrimidine-5-carbonitrile (14). Yield 83.1%. Yellow solid. Mp: 164–165 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 7.92 (d, J = 8.2 Hz, 2H, Ar–H), 7.44 (d, J = 8.1 Hz, 2H, Ar–H), 4.13 (s, 3H, –CH₃), 3.56 (t, J = 20.3 Hz, 2H, –CH₂–), 3.15 (t, J = 2.5 Hz, 1H, \equiv C–H), 2.42 (s, 3H, –CH₃). ¹³C NMR (100 MHz, DMSO- d_6 , ppm): δ 172.13, 169.62, 166.39, 145.54, 142.88, 131.32, 129.49, 129.01, 114.82, 98.19, 78.87, 73.46, 34.76, 21.09, 19.05. HR–MS (ESI) calcd for C₁₇H₁₃N₇NaS₂: [M + Na] + m/z, 402.0572; found, 402.0573.

4-(4-Isopropylphenyl)-6-((1-methyl-1H-tetrazol-5-yl)thio)-2-(prop-2-yn-1-ylthio)pyrimidine-5-carbonitrile (15). Yield 83.4%. White solid. Mp: 141–142 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 7.95 (d, J = 3.9 Hz, 2H, Ar–H), 7.50 (d, J = 8.0 Hz, 2H, Ar–H), 4.14 (d, J = 4.0 Hz, 3H, -CH $_3$), 3.60 (s, 2H, -CH $_2$ –), 3.15 (t, J = 2.5 Hz, 1H, \equiv C–H), 3.05–2.96 (m, 1H, -CH), 1.25 (d, J = 6.8 Hz, 6H, -CH $_3$). 13 C NMR (100 MHz, DMSO- d_6 , ppm): δ 172.16, 169.65, 166.36, 153.39, 145.54, 131.70, 129.19, 126.92, 114.85, 98.17, 78.89, 73.46, 34.77, 33.47, 23.47, 19.06. HR–MS (ESI) calcd for C $_{19}$ H $_{17}$ N $_7$ NaS $_2$: [M + Na] $^+$ m/z, 430.0885; found, 430.0886.

4-(3-Methoxyphenyl)-6-((1-methyl-1H-tetrazol-5-yl)thio)-2-(prop-2-yn-1-ylthio)pyrimidine-5-carbonitrile (16). Yield 89.9%. White solid. Mp: 189–192 °C. ¹H NMR (400 MHz, DMSO-d₆,

ppm): δ 7.56 (s, 1H, Ar–H), 7.54 (d, J = 3.7 Hz, 2H, Ar–H), 7.25 (dt, J = 6.9, 2.2 Hz, 1H, Ar–H), 4.12 (s, 3H, –CH₃), 3.84 (s, 3H, –CH₃), 3.61 (d, J = 2.4 Hz, 2H, –CH₂–), 3.17 (t, J = 2.5 Hz, 1H, \equiv C–H). ¹³C NMR (100 MHz, DMSO- d_6 , ppm): δ 172.22, 169.63, 166.35, 159.27, 145.52, 135.41, 130.15, 121.20, 118.09, 114.67, 114.17, 98.78, 78.95, 73.42, 55.43, 34.76, 19.11. HR–MS (ESI) calcd for $C_{17}H_{13}N_7ONaS_2$: [M + Na]⁺ m/z, 418.0521; found, 418.0521.

4-(3,4-Dichlorophenyl)-6-((1-methyl-1H-tetrazol-5-yl)thio)-2-(prop-2-yn-1-ylthio)pyrimidine-5-carbonitrile (17). Yield 89.8%. White solid. Mp: 151–152 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 8.21 (d, J = 1.8 Hz, 1H, Ar–H), 7.97 (dd, J = 8.4, 1.9 Hz, 1H, Ar–H), 7.93 (d, J = 8.4 Hz, 1H, Ar–H), 4.13 (s, 3H, –CH₃), 3.61 (d, J = 2.3 Hz, 2H, –CH₂–), 3.16 (t, J = 2.4 Hz, 1H, \equiv C–H). 13 C NMR (100 MHz, DMSO- d_6 , ppm): δ 172.45, 169.65, 164.28, 145.42, 135.21, 134.56, 131.82, 131.26, 130.80, 129.05, 114.32, 99.14, 78.77, 73.52, 34.78, 19.18. HR–MS (ESI) calcd for C₁₆H₉Cl₂N₇NaS₂: [M + Na]⁺ m/z, 455.9636; found, 455.9637.

4-((1-Methyl-1H-tetrazol-5-yl)thio)-6-phenyl-2-(prop-2-yn-1-ylthio)pyrimidine-5-carbonitrile (18). Yield 81.9%. White solid. Mp: 130–131 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 8.17 (d, J = 7.5 Hz, 2H, Ar–H), 8.07 (s, 1H, Ar–H), 7.59–7.55 (m, 2H, Ar–H), 4.10 (s, 3H, –CH₃), 3.67 (d, 2H, –CH₂–), 3.11 (t, J = 2.5 Hz,1H, \equiv C–H). ¹³C NMR (100 MHz, DMSO- d_6 , ppm): δ 172.30, 169.69, 165.47, 145.47, 137.36, 132.92, 130.85, 129.06, 114.55, 98.72, 78.78, 73.51, 34.78, 19.10. HR–MS (ESI) calcd for C₁₆H₁₁N₇NaS₂: [M + Na] + m/z, 388.0415; found, 388.0415.

4-((1-Methyl-1H-tetrazol-5-yl)amino)-2-(prop-2-yn-1-ylthio)-6-(3,4,5-trimethoxyphenyl)pyrimidine-5-carbonitrile (19). Yield 94.1%. Yellow solid. Mp: 205–206 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 7.38 (s, 2H, Ar–H), 3.93 (s, 3H, –CH₃), 3.87 (s, 6H, –CH₃), 3.85 (d, J = 1.8 Hz, 2H, –CH₂–), 3.78 (s, 3H, –CH₃), 3.16 (t, J = 2.5 Hz, 1H, \equiv C–H). ¹³C NMR (100 MHz, DMSO- d_6 , ppm): δ 171.96, 167.33, 161.41, 152.71, 150.04, 140.43, 130.11, 115.42, 106.68, 86.40, 79.95, 72.99, 60.22, 56.11, 33.61, 19.17. HR–MS (ESI) calcd for C₁₉H₁₈N₈NaO₃S: [M + Na]⁺ m/z, 461.1120; found, 461.1122.

4-(*Piperazin-1-yl*)-2-(*prop-2-yn-1-ylthio*)-6-(3,4,5-trimethoxyphenyl)pyrimidine-5-carbonitrile (20). Yield 89.0%. White solid. Mp: 242–243 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 9.72 (s, 1H, NH, D₂O exchangeable), 7.30 (s, 2H, Ar–H), 4.19 (s, 4H, -CH₂-), 4.03 (d, J = 2.1 Hz, 2H, -CH₂-), 3.86 (s, 6H, -CH₃), 3.77 (s, 3H, -CH₃), 3.34 (s, 4H, -CH₂-), 3.21 (t, J = 2.3 Hz, 1H, \equiv C-H). ¹³C NMR (100 MHz, DMSO- d_6 , ppm): δ 170.58, 169.15, 161.75, 152.60, 140.23, 130.88, 117.90, 107.18, 83.78, 80.45, 73.00, 60.18, 56.16, 45.75, 19.26. HR–MS (ESI) calcd for C₂₁H₂₄N₅O₃S: [M + H]+ m/z, 426.1600; found, 426.1602.

2-(Prop-2-yn-1-ylthio)-4-thiomorpholino-6-(3,4,5-trimethoxyphenyl)pyrimidine-5-carbonitrile (21). Yield 50.9%. Yellow solid. Mp: 152–153 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 7.28 (s, 2H, Ar–H), 4.19 (m, 4H, –CH₂–), 3.98 (d, J = 2.4 Hz, 2H, –CH₂–), 3.85 (s, 6H, –CH₃), 3.77 (s, 3H, –CH₃), 3.19 (t, J = 2.3 Hz, 1H, \equiv C–H), 2.86–2.77 (m, 4H, –CH₂–). ¹³C NMR (100 MHz, DMSO- d_6 , ppm): δ 170.71, 169.42, 161.97, 152.56, 140.21, 130.93, 117.92, 107.23, 84.24, 80.46, 72.89, 60.17, 56.17, 49.70, 26.50, 19.22. HR–MS (ESI) calcd for C₂₁H₂₂N₄O₃NaS₂: [M + Na]⁺ m/z, 465.1031; found, 465.1032.

4-Morpholino-2-(prop-2-yn-1-ylthio)-6-(3,4,5-trimethoxyphenyl)pyrimidine-5-carbonitrile (22). Yield 89.5%. White solid. Mp: 163–164 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 7.27 (s, 2H, Ar–H), 3.98 (d, J=2.5 Hz, 2H, $-CH_2-$), 3.97–3.93 (m, 4H, $-CH_2-$), 3.85 (s, 6H, $-CH_3$), 3.76 (s, 3H, $-CH_3$), 3.74 (m, 4H, $-CH_2-$), 3.19 (t, J=2.5 Hz, 1H, \equiv C-H). ¹³C NMR (100 MHz, DMSO- d_6 , ppm): δ 170.70, 169.34, 161.84, 152.58, 140.24, 130.87, 117.86, 107.20, 83.94, 80.40, 72.91, 65.91, 60.17, 56.16, 47.17, 19.23. HR–MS (ESI) calcd for $C_{21}H_{23}N_4O_4S$: [M + H]⁺ m/z, 427.1440; found, 427.1441.

4-(*Piperidin-1-yl*)-2-(*prop-2-yn-1-ylthio*)-6-(3,4,5-trimethoxyphenyl)pyrimidine-5-carbonitrile (23). Yield 90.0%. White solid. Mp: 120–121 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 7.27 (s, 2H, Ar–H), 3.97 (d, J = 2.5 Hz, 2H, -CH₂-), 3.90

(m, 4H, $-\text{CH}_2-$), 3.86 (s, 6H, $-\text{CH}_3$), 3.77 (s, 3H, $-\text{CH}_3$), 3.17 (t, J=2.5 Hz, 1H, \equiv C-H), 1.68 (s, 6H, $-\text{CH}_2-$). ^{13}C NMR (100 MHz, DMSO- d_6 , ppm): δ 170.55, 169.37, 161.46, 152.54, 140.16, 131.04, 118.01, 107.19, 83.42, 80.48, 72.81, 60.15, 56.16, 47.99, 25.58, 23.64, 19.16. HR-MS (ESI) calcd for $\text{C}_{22}\text{H}_{25}\text{N}_4\text{O}_3\text{S}$: [M + H]⁺ m/z, 425.1647; found, 425.1648.

4-((1-(2-(Dimethylamino)ethyl)-1H-tetrazol-5-yl)thio)-2-(prop-2-yn-1-ylthio)-6-(3,4,5-trimethoxyphenyl)pyrimidine-5-carbonitrile (**26**). Yield 75.9%. Yellow solid. Mp: 197–198 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 7.42 (s, 2H, Ar–H), 4.99 (t, J = 6.6 Hz, 2H, -CH₂–), 3.87 (s, 6H, -CH₃), 3.79 (s, 3H, -CH₃), 3.72 (d, J = 2.3 Hz, 2H, -CH₂–), 3.69 (t, J = 6.6 Hz, 2H, -CH₂–), 3.25 (t, J = 2.5 Hz, 1H, \equiv C–H), 2.83 (s, 6H, -CH₃). 13 C NMR (100 MHz, DMSO- d_6 , ppm): δ 171.96, 169.87, 165.62, 152.90, 146.30, 141.06, 129.01, 115.03, 106.80, 98.30, 79.26, 73.45, 60.29, 56.19, 54.14, 42.69, 42.38, 19.36. HR–MS (ESI) calcd for $C_{22}H_{25}N_8O_3S_2$: [M + H]⁺ m/z, 513.1491; found, 513.1491.

4-((1H-1,2,4-Triazol-3-yl)thio)-2-(prop-2-yn-1-ylthio)-6-(3,4,5-trimethoxyphenyl)pyrimidine-5-carbonitrile (27). Yield 80.9%. White solid. Mp: 234–235 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 8.85 (s, 1H, NH, D₂O exchangeable), 7.41 (s, 2H, Ar–H), 3.87 (s, 6H, –CH₃), 3.79 (s, 3H, –CH₃), 3.78–3.78 (s, 2H, –CH₂–), 3.15 (t, J = 2.4 Hz, 1H, \equiv C–H). ¹³C NMR (100 MHz, DMSO- d_6 , ppm): δ 172.22, 169.63, 166.35, 159.27, 145.52, 135.41, 130.15, 121.20, 118.09, 114.67, 114.17, 98.78, 78.95, 73.42, 55.43, 34.76, 19.11. HR–MS (ESI) calcd for C₁₉H₁₆N₆NaO₃S₂: [M + Na]⁺ m/z, 463.0623; found, 463.0622.

4-((1-Methyl-1H-imidazol-2-yl)thio)-2-(prop-2-yn-1-ylthio)-6-(3,4,5-trimethoxyphenyl)pyrimidine-5-carbonitrile (28). Yield 32.8%. White solid. Mp: 176–177 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 7.62 (s, 1H, -CH=), 7.38 (s, 2H, Ar-H), 7.22 (s, 1H, -CH=), 3.87 (s, 6H, -CH₃), 3.79 (s, 3H, -CH₃), 3.68 (s, 2H, -CH₂-), 3.67 (s, 3H, -CH₃), 3.20 (t, J = 2.2 Hz, 1H, \equiv C-H). ¹³C NMR (100 MHz, DMSO- d_6 , ppm): δ 173.22, 171.82, 165.81, 152.83, 140.84, 130.39, 130.16, 129.27, 126.44, 115.07, 106.86, 97.48, 79.44, 73.36, 60.25, 56.18, 33.68, 19.22. HR-MS (ESI) calcd for $C_{21}H_{19}N_5NaO_3S_2$: [M + Na] + m/z, 476.0827; found, 476.0828.

2-(*Prop-2-yn-1-ylthio*)-4-(*pyridin-2-ylthio*)-6-(3,4,5-trimethoxyphenyl)pyrimidine-5-carbonitrile (29). Yield 80.2%. Yellow solid. Mp: 180–181 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 8.76–8.58 (m, 1H, Ar–H), 7.97 (td, J = 7.7, 1.7 Hz, 1H, Ar–H), 7.86 (d, J = 7.8 Hz, 1H, Ar–H), 7.55 (dt, J = 15.7, 7.8 Hz, 1H, Ar–H), 7.38 (s, 2H, Ar–H), 3.86 (s, 6H, –CH₃), 3.78 (s, 3H, –CH₃), 3.71 (d, J = 2.2 Hz, 2H, –CH₂–), 3.14 (t, J = 2.3 Hz, 1H). 13 C NMR (100 MHz, DMSO- d_6 , ppm): δ 172.90, 171.53, 165.73, 152.80, 150.76, 149.32, 140.76, 138.08, 131.26, 129.36, 124.71, 115.22, 106.82, 98.15, 79.54, 73.35, 60.24, 56.14, 19.20. HR–MS (ESI) calcd for C₂₂H₁₈N₄NaO₃S₂: [M + Na]⁺ m/z, 473.0718; found, 473.0715.

4-((5-Aminothiophen-2-yl)thio)-2-(prop-2-yn-1-ylthio)-6-(3,4,5-trimethoxyphenyl)pyrimidine-5-carbonitrile (30). Yield 89.3%. Yellow solid. Mp: 189–190 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 7.87 (s, 2H, -CH=), 7.42 (s, 2H, Ar-H), 3.95 (d, J = 2.1 Hz, 2H, -CH₂-), 3.86 (s, 6H, -CH₃), 3.79 (s, 3H, -CH₃), 3.17 (d, J = 2.2 Hz, 1H, \equiv C-H). 13 C NMR (100 MHz, DMSO- d_6 , ppm): δ 173.78, 171.83, 171.36, 165.58, 152.84, 140.98, 129.19, 114.91, 106.95, 97.68, 79.43, 73.44, 60.27, 56.19, 19.55. HR-MS (ESI) calcd for $C_{21}H_{18}N_4NaO_3S_3$: $[M+Na]^+$ m/z, 493.0439; found, 493.0438.

4-((5-Methyl-1,3,4-thiadiazol-2-yl)thio)-2-(prop-2-yn-1-ylthio)-6-(3,4,5-trimethoxyphenyl)pyrimidine-5-carbonitrile (31). Yield 76.5%. White solid. Mp: 186–187 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 7.44 (s, 2H, Ar–H), 3.93 (d, J=2.4 Hz, 2H, –CH₂–), 3.87 (s, 6H, –CH₃), 3.80 (s, 3H, –CH₃), 3.20 (t, J=2.5 Hz, 1H, \equiv C–H), 2.86 (s, 1H, –CH₃). ¹³C NMR (100 MHz, DMSO- d_6 , ppm): δ 171.88, 171.26, 169.59, 165.64, 154.06, 152.86, 141.07, 129.09, 114.85, 106.96, 98.14, 79.53, 73.43, 60.27, 56.19, 19.54, 15.53. HR–MS (ESI) calcd for C₂₀H₁₇N₅NaO₃S₃: [M + Na] + m/z, 494.0391; found, 494.0389.

4-((5-Amino-1,3,4-thiadiazol-2-yl)thio)-2-(prop-2-yn-1-ylthio)-6-(3,4,5-trimethoxyphenyl)pyrimidine-5-carbonitrile (32). Yield 65.1%. White solid. Mp: 181–182 °C. ¹H NMR (400 MHz,

DMSO- d_6 , ppm): δ 7.83 (s, 2H, NH, D₂O exchangeable), 7.43 (d, J = 0.6 Hz, 2H, Ar–H), 3.94 (t, J = 7.4 Hz, 2H, -CH₂–), 3.87 (s, 6H, -CH₃), 3.79 (d, J = 0.8 Hz, 3H, -CH₃), 3.18 (dd, J = 2.5, 1.6 Hz, 1H, \equiv C–H). 13 C NMR (100 MHz, DMSO- d_6 , ppm): δ 173.82, 171.81, 171.41, 165.59, 152.84, 140.96, 138.79, 129.21, 114.92, 106.95, 97.71, 79.44, 73.48, 60.26, 56.19, 19.55.

4-((1,3,4-Thiadiazol-2-yl)thio)-2-(prop-2-yn-1-ylthio)-6-(3,4,5-trimethoxyphenyl)pyrimidine-5-carbonitrile (33). Yield 81.1%. White solid. Mp: 228–229 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 9.96 (s, 1H, -CH=), 7.45 (s, 2H, Ar-H), 3.94 (d, J = 2.5 Hz, 2H, -CH₂-), 3.87 (s, 6H, -CH₃), 3.79 (s, 3H, -CH₃), 3.20 (t, J = 2.5 Hz, 1H, \equiv C-H). ¹³C NMR (100 MHz, DMSO- d_6 , ppm): δ 171.83, 169.14, 165.64, 159.31, 155.09, 152.88, 141.10, 129.10, 114.85, 106.99, 98.24, 79.51, 73.53, 60.28, 56.20, 19.57.

2-(*Prop-2-yn-1-ylthio*)-4-(*thiazol-2-ylthio*)-6-(3,4,5-trimethoxyphenyl)pyrimidine-5-carbonitrile (34). Yield 81.1%. White solid. Mp: 172–173 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): 8.20 (d, J=3.2 Hz, 1H, -CH \Longrightarrow), 8.11 (d, J=3.2 Hz, 1H, -CH \Longrightarrow), 7.42 (s, 2H, Ar \Longrightarrow H, 3.87 (s, 6H, \ast CH₃), 3.85 (s, 2H, \ast CH₂ \Longrightarrow), 3.79 (s, 3H, \ast CH₃), 3.18 (s, 1H, \ast C \Longrightarrow C \Longrightarrow CNMR (100 MHz, DMSO- d_6 , ppm): δ 171.75, 171.21, 165.61, 152.84, 150.61, 144.13, 140.94, 129.20, 127.64, 114.88, 106.93, 97.72, 79.49, 79.11, 73.49, 60.26, 56.17, 19.41. HR \ast MS (ESI) calcd for C₂₀H₁₆N₄NaO₃S₃: [M + Na]* m/z, 479.0282; found, 479.0284.

4-((4-Methylthiazol-2-yl)thio)-2-(prop-2-yn-1-ylthio)-6-(3,4,5-trimethoxyphenyl)pyrimidine-5-carbonitrile (35). Yield 76.7%. White solid. Mp: 226–227 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 7.75 (d, J = 0.9 Hz, 1H, -CH=), 7.41 (s, 2H, Ar=H), 3.86 (s, 6H, -CH₃), 3.86 (d, J = 2.8 Hz, 2H, -CH₂-), 3.79 (s, 3H, -CH₃), 3.18 (t, J = 2.6 Hz, 1H), 2.46 (d, J = 0.8 Hz, 3H, -CH₃). ¹³C NMR (100 MHz, DMSO- d_6 , ppm): δ 171.72, 171.36, 165.67, 153.78, 152.84, 149.35, 140.94, 129.23, 121.97, 114.88, 106.95, 97.72, 79.51, 73.41, 60.26, 56.19, 19.38, 16.73. HR=MS (ESI) calcd for $C_{21}H_{18}N_4NaO_3S_3$: [M + Na]* m/z, 493.0439; found, 493.0434.

4-((4,5-Dimethylthiazol-2-yl)thio)-2-(prop-2-yn-1-ylthio)-6-(3,4,5-trimethoxyphenyl)pyrimidine-5-carbonitrile (36). Yield 82.9%. White solid. Mp: 192–193 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 7.23 (s, 2H, Ar–H), 3.70 (s, 6H, –CH₃), 3.62 (s, 3H, –CH₃), 3.22 (s, 2H, –CH₂–), 3.02 (s, 1H, \equiv C–H), 2.29 (s, 3H, –CH₃), 2.19 (s, 3H, –CH₃). ¹³C NMR (100 MHz, DMSO- d_6 , ppm): δ 171.72, 171.36, 165.67, 153.78, 152.84, 149.35, 140.94, 129.23, 121.97, 114.88, 106.95, 97.72, 79.51, 73.41, 60.26, 56.19, 19.38, 16.73, 19.24. HR–MS (ESI) calcd for C₂₂H₂₁N₄O₃S₃: [M + H]+ m/z, 485.0776; found, 485.0774.

4-(Benzo[d]thiazol-2-ylthio)-2-(prop-2-yn-1-ylthio)-6-(3,4,5-trimethoxyphenyl)pyrimidine-5-carbonitrile (37). Yield 69.0%. White solid. Mp: 223–224 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 8.25–8.17 (m, 1H, Ar–H), 8.10 (d, J = 7.6 Hz, 1H, Ar–H), 7.58 (dtd, J = 16.4, 7.3, 1.3 Hz, 2H, Ar–H), 7.47 (s, 2H, Ar–H), 3.92 (d, J = 2.4 Hz, 2H, $-CH_2$ -), 3.88 (s, 6H, $-CH_3$), 3.80 (s, 3H, $-CH_3$), 3.15 (t, J = 2.5 Hz, 1H, \equiv C–H). 13 C NMR (100 MHz, DMSO- d_6 , ppm): δ 171.72, 169.72, 165.59, 155.24, 152.86, 151.49, 141.09, 136.73, 129.15, 126.81, 126.23, 122.92, 122.16, 114.86, 107.01, 98.14, 79.40, 73.39, 60.27, 56.20, 19.52. HR–MS (ESI) calcd for $C_{24}H_{18}N_4NaO_3S_3$: $[M+Na]^+$ m/z, 529.0439; found, 529.0439.

4-(3-Chlorophenyl)-2-(prop-2-yn-1-ylthio)-6-(thiazol-2-ylthio)-pyrimidine-5-carbonitrile (38). Yield 65.4%. White solid. Mp: 171–172 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 8.24–8.21 (m, 1H, Ar–H), 8.13 (d, J = 2.9 Hz, 1H, Ar–H), 8.03 (s, 1H, Ar–H), 7.96 (d, J = 7.6 Hz, 1H, Ar–H), 7.75 (d, J = 8.0 Hz, 1H, -CH=), 7.66 (t, J = 7.8 Hz, 1H, -CH=), 3.80 (s, 2H, -CH₂-), 3.15 (s, 1H, =C-H). ¹³C NMR (100 MHz, DMSO- d_6 , ppm): δ 172.10, 171.43, 165.05, 150.21, 144.34, 136.33, 133.49, 131.82, 130.75, 128.75, 127.96, 127.72, 114.35, 98.60, 79.11, 73.82, 19.31. HR-MS (ESI) calcd for C₁₇H₉ClN₄NaS₃: [M + Na]⁺ m/z, 422.9576; found, 422.9577.

4-(4-Chlorophenyl)-2-(prop-2-yn-1-ylthio)-6-(thiazol-2-ylthio)-pyrimidine-5-carbonitrile (**39**). Yield 73.2%. White solid. Mp: 136–137 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 8.21 (t, J = 10.7 Hz, 1H, –CH=), 8.13 (d, J = 3.3 Hz, 1H, –CH=), 8.03 (d, J = 8.6 Hz, 2H, Ar–H), 7.71 (d, J = 8.6 Hz, 2H, Ar–H), 3.79 (d, J = 2.4 Hz, 2H,

 $-\text{CH}_2-$), 3.15 (t, J=2.5 Hz, 1H, \equiv C-H). ¹³C NMR (100 MHz, DMSO- d_6 , ppm): δ 172.00, 171.47, 165.31, 150.28, 144.31, 137.20, 133.13, 130.92, 128.97, 127.92, 114.47, 98.23, 79.08, 73.86, 19.27. HR-MS (ESI) calcd for $\text{C}_{17}\text{H}_9\text{ClN}_4\text{NaS}_3$: [M + Na]⁺ m/z, 422.9576; found, 422.9578.

4-(4-Bromophenyl)-2-(prop-2-yn-1-ylthio)-6-(thiazol-2-ylthio)-pyrimidine-5-carbonitrile (40). Yield 78.1%. White solid. Mp: 238–239 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 8.23 (d, J = 3.3 Hz, 1H, -CH=), 8.13 (d, J = 3.3 Hz, 1H, -CH=), 7.97–7.91 (m, 2H, Ar–H), 7.88–7.82 (m, 2H, Ar–H), 3.79 (d, J = 2.6 Hz, 2H, -CH₂–), 3.14 (t, J = 2.6 Hz, 1H, \equiv C–H). 13 C NMR (100 MHz, DMSO- d_6 , ppm): δ 172.03, 171.48, 165.43, 150.29, 144.31, 133.50, 131.91, 131.04, 127.91, 126.18, 114.45, 98.20, 79.06, 73.86, 19.29. HR–MS (ESI) calcd for C₁₇H₉BrN₄NaS₃: [M + Na]⁺ m/z, 466.9070; found, 466.9073.

4-(3,4-Dichlorophenyl)-2-(prop-2-yn-1-ylthio)-6-(thiazol-2-ylthio)pyrimidine-5-carbonitrile (41). Yield 88.0%. White solid. Mp: 142–143 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 8.23 (d, J = 3.1 Hz, 2H, Ar–H), 8.13 (d, J = 3.3 Hz, 1H, Ar–H), 7.98 (dd, J = 8.4, 1.7 Hz, 1H, –CH=), 7.91 (d, J = 8.4 Hz, 1H, –CH=), 3.80 (d, J = 2.3 Hz, 2H, –CH₂–), 3.15 (s, 1H, \equiv C–H). ¹³C NMR (100 MHz, DMSO- d_6 , ppm): δ 172.14, 171.46, 164.05, 150.12, 144.36, 135.06, 134.74, 131.74, 131.14, 130.89, 129.11, 127.99, 114.26, 98.58, 79.06, 73.84, 19.35. HR–MS (ESI) calcd for C₁₇H₈C₁₂N₄NaS₃: [M + Na]⁺ m/z, 456.9186; found, 456.9187.

2-(*Prop-2-yn-1-ylthio*)-4-(*thiazol-2-ylthio*)-6-(*p-tolyl*)*pyrimidine-5-carbonitrile* (*42*). Yield 72.8%. White solid. Mp: 156–157 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 8.21 (d, J=3.3 Hz, 1H, –CH=), 8.12 (d, J=3.3 Hz, 1H, –CH=), 7.93 (d, J=8.1 Hz, 2H, Ar–H), 7.43 (d, J=8.1 Hz, 2H, Ar–H), 3.80 (d, J=2.5 Hz, 2H, –CH₂–), 3.15 (t, J=2.4 Hz, 1H, \equiv C–H), 2.41 (d, J=6.0 Hz, 3H, –CH₃). ¹³C NMR (100 MHz, DMSO- d_6 , ppm): δ 172.22, 169.63, 166.35, 159.27, 145.52, 135.41, 130.15, 121.20, 118.09, 114.67, 114.17, 98.78, 78.95, 73.42, 55.43, 34.76, 19.11. HR–MS (ESI) calcd for $C_{18}H_{12}N_4NaS_3$: [M + Na] * m/z, 403.0122; found, 403.0123.

4-(2-Oxo-2H-chromen-6-yl)-2-(prop-2-yn-1-ylthio)-6-(thiazol-2-ylthio)pyrimidine-5-carbonitrile (43). Yield 90.8%. Yellow solid. Mp: 226–227 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 8.37 (d, J = 1.9 Hz, 1H, Ar–H), 8.23 (d, J = 3.3 Hz, 1H, Ar–H), 8.22 (d, J = 2.5 Hz, 1H, -CH=), 8.20 (s, 1H, -CH=), 8.13 (d, J = 3.3 Hz, 1H, -CH=), 7.64 (d, J = 8.7 Hz, 1H, -CH=), 6.62 (d, J = 9.6 Hz, 1H, Ar–H), 3.80 (d, J = 2.4 Hz, 2H, -CH₂–), 3.16 (t, J = 2.4 Hz, 1H, = C–H). 13 C NMR (100 MHz, DMSO- d_6 , ppm): δ 172.02, 171.52, 165.02, 159.35, 155.67, 150.23, 144.34, 143.75, 132.30, 130.47, 129.70, 127.97, 118.99, 117.34, 117.01, 114.53, 98.15, 79.08, 73.94, 19.32. HR–MS (ESI) calcd for C₂₀H₁₀N₄O₂NaS₃: [M + Na]⁺ m/z, 456.9864; found, 456.9862.

2-(*Prop-2-yn-1-ylthio*)-4-(*pyridin-4-yl*)-6-(*thiazol-2-ylthio*)-pyrimidine-5-carbonitrile (44). Yield 76.3%. White solid. Mp: 189–190 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 8.87 (dd, J = 4.5, 1.5 Hz, 2H, Ar–H), 8.24 (d, J = 3.3 Hz, 1H, -CH=), 8.14 (d, J = 3.3 Hz, 1H, -CH=), 7.91 (dd, J = 4.5, 1.6 Hz, 2H, Ar–H), 3.79 (d, J = 2.5 Hz, 2H, -CH₂-), 3.14 (t, J = 2.6 Hz, 1H, \equiv C–H). ¹³C NMR (100 MHz, DMSO- d_6 , ppm): δ 190.70, 176.34, 166.45, 165.53, 149.87, 142.64, 140.34, 121.31, 118.64, 117.01, 96.91, 76.90, 73.97, 25.02. HR–MS (ESI) calcd for C₁₆H₉N₃NaS₃: [M + Na]⁺ m/z, 389.9918; found, 389.9919.

2-(*Prop-2-yn-1-ylthio*)-4-(1*H-pyrrol-2-yl*)-6-(thiazol-2-ylthio)-pyrimidine-5-carbonitrile (45). Yield 80.1%. White solid. Mp: 255–256 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 12.05 (s, 1H, NH, D₂O exchangeable), 8.17 (d, J = 3.3 Hz, 1H, -CH=), 8.08 (d, J = 3.3 Hz, 1H, -CH=), 7.30 (s, 1H, -CH=), 6.42 (dt, J = 4.2, 2.3 Hz, 1H, -CH=), 3.94 (d, J = 2.5 Hz, 2H, -CH₂-), 3.15 (t, J = 2.6 Hz, 1H, =C-H). 13 C NMR (100 MHz, DMSO- d_6 , ppm): δ 171.25, 170.38, 155.31, 150.94, 143.92, 127.38, 127.03, 125.56, 116.89, 115.36, 112.01, 91.23, 79.30, 73.98, 19.04. HR-MS (ESI) calcd for C₁₅H₉N₅NaS₃: [M + Na]⁺ m/z, 377.9918; found, 377.9917.

4-(4-Chlorophenyl)-6-((1-(2-(dimethylamino)ethyl)-1H-tetrazol-5-yl)thio)-2-(prop-2-yn-1-ylthio)pyrimidine-5-carbonitrile (46).

Yield 89.0%. White solid. Mp: 194–196 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 8.00 (d, J = 8.5 Hz, 2H, Ar–H), 7.69 (d, J = 8.5 Hz, 2H, Ar–H), 4.94 (t, J = 6.1 Hz, 2H, -CH₂–), 3.61 (s, 4H, -CH₂–), 3.10 (s, 1H, \equiv C–H), 2.85 (s, 6H, -CH₃). ¹³C NMR (100 MHz, DMSO- d_6 , ppm): δ 172.32, 169.75, 165.42, 146.12, 137.45, 132.85, 130.83, 129.10, 114.52, 98.73, 78.84, 73.55, 54.27, 42.58, 19.16. HR–MS (ESI) calcd for C₁₉H₁₈ClN₈S₂: [M + H]⁺ m/z, 457.0784; found, 457.0785.

4-(3-Chlorophenyl)-6-((1-(2-(dimethylamino)ethyl)-1H-tetrazol-5-yl)thio)-2-(prop-2-yn-1-ylthio)pyrimidine-5-carbonitrile (47). Yield 75.2%. Yellow solid. Mp: 179–180 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 8.02 (s, 1H, Ar–H), 7.97 (d, J = 7.9 Hz, 1H, Ar–H), 7.81–7.75 (m, 1H, –CH=), 7.69 (t, J = 7.9 Hz, 1H, –CH=), 4.99 (t, J = 6.6 Hz, 2H, –CH₂–), 3.69 (t, J = 5.1 Hz, 2H, –CH₂–), 3.67 (d, J = 2.2 Hz, 2H, –CH₂–), 3.21 (t, J = 2.4 Hz, 1H, =C–H), 2.84 (s, 6H, –CH₃). ¹³C NMR (100 MHz, DMSO- d_6 , ppm): δ 172.34, 169.87, 165.06, 146.17, 136.11, 133.61, 132.04, 130.93, 128.64, 127.64, 114.45, 99.20, 78.85, 73.69, 54.14, 42.72, 42.41, 19.24. HR–MS (ESI) calcd for C₁₉H₁₈ClN₈S₂: [M + H]⁺ m/z, 457.0784; found, 457.0785.

4-((1-(2-(Dimethylamino)ethyl)-1H-tetrazol-5-yl)thio)-2-(prop-2-yn-1-ylthio)-6-(p-tolyl)pyrimidine-5-carbonitrile (48). Yield 46.0%. White solid. Mp: 190–191 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 7.94 (d, J = 8.2 Hz, 2H, Ar–H), 7.45 (d, J = 8.2 Hz, 2H, Ar–H), 4.97 (t, J = 6.4 Hz, 2H, -CH₂-), 3.65 (d, J = 2.3 Hz, 4H, -CH₂-), 3.20 (t, J = 2.3 Hz, 1H, \equiv C-H), 2.81 (s, 6H, -CH₃), 2.43 (s, 3H, -CH₃). 13 C NMR (100 MHz, DMSO- d_6 , ppm): δ 172.05, 169.89, 166.23, 146.24, 142.93, 131.32, 129.54, 129.01, 114.87, 98.23, 78.92, 73.67, 54.29, 42.83, 42.49, 21.11, 19.13. HR–MS (ESI) calcd for C_{20} H₂₁N₈S₂: [M + H]⁺ m/z, 437.1331; found, 437.1330.

4-(3,4-Dichlorophenyl)-6-((1-(2-(dimethylamino)ethyl)-1H-tetrazol-5-yl)thio)-2-(prop-2-yn-1-ylthio)pyrimidine-5-carbonitrile (49). Yield 46.0%. White solid. Mp: 174–175 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 8.23 (s, 1H, Ar–H), 7.97 (dd, J = 17.7, 8.2 Hz, 2H, Ar–H), 4.98 (t, J = 6.3 Hz, 2H, -CH₂–), 3.69 (s, 2H, -CH₂–), 3.68 (s, 2H, -CH₂–), 3.21 (s, 1H, \equiv C–H), 2.83 (s, 3H, -CH₃). ¹³C NMR (100 MHz, DMSO- d_6 , ppm): δ 172.40, 169.87, 164.10, 146.12, 135.29, 134.53, 131.88, 131.33, 130.77, 129.03, 114.35, 99.21, 78.82, 73.70, 54.15, 42.74, 42.44, 19.27. HR-MS (ESI) calcd for C₁₉H₁₇C₁₂N₈S₂: [M + H]⁺ m/z, 491.0395; found, 491.0395.

4-((1-(2-(Dimethylamino)ethyl)-1H-tetrazol-5-yl)thio)-6-(2-oxo-2H-chromen-6-yl)-2-(prop-2-yn-1-ylthio)pyrimidine-5-carbonitrile (50). Yield 72.4%. Yellow solid. Mp: 197–198 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 8.39 (s, 1H, Ar–H), 8.27–8.19 (m, 2H, Ar–H), 7.66 (d, J = 8.7 Hz, 1H, Ar–H), 6.63 (d, J = 9.6 Hz, 1H, Ar–H), 5.00 (t, J = 6.5 Hz, 2H, -CH₂-), 3.69 (d, J = 4.9 Hz, 2H, -CH₂-), 3.66 (d, J = 1.9 Hz, 2H, -CH₂-), 3.23 (t, J = 2.3 Hz, 1H, \equiv C–H), 2.83 (s, 6H, -CH₃). 13 C NMR (100 MHz, DMSO- d_6 , ppm) δ 172.25, 169.97, 165.07, 159.32, 155.78, 146.19, 143.74, 132.17, 130.26, 129.65, 119.12, 117.43, 117.17, 114.64, 98.72, 78.82, 73.79, 54.23, 42.81, 42.46, 42.34, 19.22. HR–MS (ESI) calcd for $C_{22}H_{18}N_8NaO_2S_3$: [M + Na] $^+$ m/z, 513.0892; found, 513.0893.

2-(*Allylthio*)-4-(4-chlorophenyl)-6-(thiazol-2-ylthio)pyrimidine-5-carbonitrile (*51*). Yield 56.0%. White solid. Mp: 109–110 °C. ¹H NMR (400 MHz, CDCl₃): δ 8.19 (d, J = 3.3 Hz, 1H, Ar–H), 8.11 (d, J = 3.3 Hz, 1H, Ar–H), 8.03–7.85 (m, 2H, Ar–H), 7.77–7.60 (m, 2H, Ar–H), 5.70 (dd, J = 16.9, 10.1 Hz, 1H, –CH=), 5.28–4.98 (m, 2H, –CH₂–), 3.56 (d, J = 6.8 Hz, 2H, –CH₂–). ¹³C NMR (101 MHz, DMSO): δ 173.13, 171.37, 165.31, 150.35, 144.44, 137.05, 133.23, 132.71, 130.83, 128.94, 128.09, 118.36, 114.57, 97.88, 33.16. HR–MS (ESI) calcd for C₁₇H₁₁ClN₄NaS₃: [M + Na]⁺m/z, 424.9732; found, 424.9730.

4-(4-Chlorophenyl)-2-(propylthio)-6-(thiazol-2-ylthio)-pyrimidine-5-carbonitrile (52). Yield 48.2%. White solid. Mp: 112–113 °C. ¹H NMR (400 MHz, DMSO): δ 8.25 (d, J = 3.3 Hz, 1H, Ar–H), 8.15 (d, J = 3.3 Hz, 1H, Ar–H), 8.02–7.89 (m, 2H, Ar–H), 7.70 (dd, J = 8.9, 2.1 Hz, 2H, Ar–H), 2.82 (t, J = 7.2 Hz, 2H, -CH₂-), 1.46 (dd, J = 14.5, 7.3 Hz, 2H, -CH₂-), 0.86 (t, J = 7.3 Hz, 3H, -CH₃). 13 C NMR (101 MHz, DMSO): δ 173.90, 171.38, 165.19, 150.25, 144.56, 136.97, 133.29, 130.79, 128.93, 128.26, 114.65, 97.58,

32.44, 21.92, 13.05. HR–MS (ESI) calcd for $C_{17}H_{13}ClN_4NaS_3$: [M + Na] $^+m/z$, 426.9889; found, 426.9890.

2-(((1-Benzyl-1H-1,2,3-triazol-4-yl)methyl)thio)-4-(4-chlorophenyl)-6-(thiazol-2-ylthio)pyrimidine-5-carbonitrile (**54**). Yield 32.6%. White solid. Mp: 177–178 °C. ¹H NMR (400 MHz, DMSO- d_6 , ppm): δ 8.08 (d, J = 3.3 Hz, 1H, -CH=), 7.99 (d, J = 2.4 Hz, 2H, Ar-H), 7.97 (s, 1H, Ar-H), 7.87 (s, 1H, Ar-H), 7.69 (d, J = 8.6 Hz, 2H, Ar-H), 7.36 (td, J = 8.8, 4.4 Hz, 3H, Ar-H), 7.30–7.23 (m, 2H, Ar-H), 5.53 (s, 2H, S-CH₂-), 4.24 (s, 2H, -CH₂-). ¹³C NMR (100 MHz, DMSO- d_6 , ppm): δ 190.71, 176.32, 166.40, 165.5, 142.63, 134.33, 133.78, 130.78, 129.37, 128.96, 128.63, 127.62, 125.79, 122.96, 118.62, 117.05, 96.95, 57.32, 35.20. HR-MS (ESI) calcd for $C_{24}H_{16}ClN_7NaS_3$: [M + Na]+ m/z, 556.0216; found, 556.0217.

Reagents. Rabbit monoclonal antibodies against DCN1 (ab181233), Cullin1 (ab75817), Cullin2 (ab166917), Cullin3 (ab75851), UBE2M (ab109507), p21 (ab109520), and Cullin5 (ab184177) were purchased from Abcam Biotechnology (Cambridge, UK). Anti-DCN1 (GWB-E3D700) antibody was from Genway Biotech (San Diego, CA). Mouse monoclonal antibody against CyclinE1 (4129) was purchased from Cell Signaling Technologies. Rabbit polyclonal antibodies against CDT1 (14382-1-AP), HO-1/HMOX1 (10701-1-AP), NQO1 (11451-1-AP), NRF2 (16396-1-AP), CUL4A (14851-1-AP), and CUL4B (12916-1-AP) were purchased from Proteintech (Wuhan, China).

E. coli DH5 α and BL21 (DE3) were obtained from Life technologies. E. coli BL21-AI was purchased from Invitrogen. Kanamycin, isopropyl-β-D-galactopyranoside (IPTG) (I8070), L-(+)-arabinose (L8060), bicinchoninic acid (BCA) Protein Assay kit (PC0020), 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) (M8180), and glutathione were purchased from Solarbio (China). Ni-beads and GST-trap column were purchased from QIAGEN (U.S.). Annexin-V-FITC/PI apoptosis kit (KGA106) was purchased from Keygen Biotech (China). Enhanced chemiluminescence (QL228436) and Lipofectamine RNAiMAX Reagent (13778030) were purchased from Thermo Fisher Scientific. HTRF detection buffer (62SDBRDD), streptavidin-XL665 (610SAXLA), and anti-GST-cryptate gold (61GSTKLA) were purchased from Cisbio Bioassays. NEDD8 Conjugation Initiation Kit was obtained from Boston BioChem (Cambridge, MA). 5000×Sypro Orange dye was purchased from Invitrogen. Super Streptavidin (SSA) sensor was purchased from Fortebio.

Cell Lines. Cells were cultivated in corresponding medium supplemented with 10% FBS and 5% CO₂ at 37 °C. Human esophageal carcinoma cell lines EC109, EC9706, TE-1, human lung carcinoma cell lines PC9, AS49, and H1975, prostate carcinoma cell line PC3, and liver carcinoma cell lines BEL-7402, SMMC-7721, and ZIP177 were obtained from Cell Bank of the Chinese Academy of Sciences (Shanghai, China). Human esophageal carcinoma cell lines KYSE-70 and KYSE-140 and human immortalized normal esophageal epithelial cell Het-1A were obtained as gifts from the First Affiliated Hospital of Zhengzhou University, which were purchased from the American Type Culture Collection (Manassas, VA). Human gastric epithelial mucosa cell line GES-1 and human breast cancer cell line MCF-7 were purchased from the State Key Laboratory of Molecular Oncology, Chinese Academy of Medical Sciences (Beijing, China).

Expression and Purification of DCN1–5 Proteins. Human pDEST17-DCN1 (residues 58–259) plasmid containing N-terminal His₆ tag was a gift from Dr. Shaomeng Wang's group at the University of Michigan, Ann Arbor. DCN1 (residues 58–259), DCN2 (residues 62–259), DCN3 (residues 86–304), DCN4 (residues 102–292), and DCN5 (residues 47–237) were cloned into an N-terminal GST tag plasmid pGEX4T-1. The plasmids pDEST17-DCN1 and pGEX4T-1-DCN1–5 were transformed into *E. coli* BL21-AI and BL21(DE3), respectively. A preculture was grown in medium containing 50 μ g/mL ampicillin at 37 °C. When the cells grew until OD₆₀₀ > 0.7 after about 5–8 h, 4 mg/mL L-(+)-Arabinose and 0.25 mM isopropyl- β -D-galactopyranoside (IPTG) were added to the culture medium to induce the protein synthesis for 8 h at 20 °C. Subsequently, the cells were collected and disrupted in ice-cold washing buffer (25 mM Tris-HCl pH 7.5, 1 mM DTT, 200 mM

NaCl) by sonication. The soluble and pellet fractions were separated by centrifugation at 12 000g for 15 min. After filtration, cleared filtrate containing N-terminal His $_6$ tag protein was applied to a Ni-beads column and washed with washing buffer containing different concentration of imidazole. The filtrate containing N-terminal GST tag protein was then purified by GST-trap column and washed with washing buffer containing 10 mM glutathione. Afterward, the molecular weights of these proteins were confirmed by Commassie Blue staining, and the concentration of the recombinant proteins was determined, using a Bicinchoninic Acid (BCA) Protein Assay kit.

Competitive FP Binding Assay. The fluorescence polarization (FP) competitive binding assays were performed similarly as described previously. 30,32 FP assays were carried out in 96-well, black, round-bottom microtiter plates. The FAM-labeled fluorescent probe (FAM-782) with a K_d value of 120.43 nM for DCN1 and the analyzing protocol³⁵ were kindly provided by Shaomeng Wang's group at the University of Michigan, Ann Arbor for compound screening.³⁰ DCN1 containing N-terminal His₆ tag was used for FP binding assay. DCN1 and the probe were mixed to a final volume of 100 μ L in the assay buffer (100 mM phosphate buffer, pH = 6.5, 0.02% Tween-20). After being shaken at room temperature for 30 min, the plates were measured by PerkinElmer Envision microplate reader at an excitation wavelength of 485 nm and an emission wavelength of 530 nm. The data were analyzed by SPSS 20 and GraphPad Prism 5 software. K_i values were calculated on the basis of the methods described earlier. The equation is $K_i = [I]_{50}/([L]_{50})$ $K_d + [P]_0/K_d + 1$), where $[I]_{50}$ is the concentration of the inhibitor at 50% inhibition, [L]₅₀ is the concentration of the ligand at 50% inhibition, $K_{\rm d}$ is the dissociation constant of the protein-ligand complex, and $[P]_0$ is the concentration of the protein at 0% inhibition.

HTRF Assay. HTRF assays were carried out in 384-well, white, round-bottom microtiter plates at a final volume of 20 μ L per well. The assay cocktail was prepared as a mixture of HTRF detection buffer (62SDBRDD), Streptavidin-XL665 (610SAXLA), Anti-GST-Cryptate Gold (61GSTKLA), 5 nM GST-DCN1-5, 5 nM AcUBE2M1-21-biotin, in assay buffer (100 mM phosphate buffer, pH = 6.5, 0.02% Tween-20). The AcUBE2M $^{1-21}$ -biotin peptide was synthesized by China Peptide Co. (Shanghai, China), and its sequence is Ac-MIKLFSLKQQKKEEESAGGTK-biotin. The cocktail was incubated for 30 min at room temperature. Compounds to be screened were added to assay plates, and the control well was added in the same volume of DMSO solution. The assay mixture was incubated for 30 min at room temperature before the measurement of the HTRF signal with a PerkinElmer Envision microplate reader equipped with modules for excitation at 320 nm and emission at 615 and 665 nm. The data were analyzed by SPSS 20 and GraphPad Prism 5 software. Ki values were calculated on the basis of the methods described earlier.

Normal Thermal Shift Assay. The assay was performed on a C1000 Touch Thermal cycler-CFX96TM Real-Time PCR (BIO-RAD), using the 96-well thin-wall PCR plate. The buffer used in this experiment is 100 mM phosphate buffer, pH = 6.5, 0.02% Tween-20. A total volume of 25 μ L of solution containing 0.2 mg/mL DCN1 protein, compounds (DC-2, DC-2N, and NAcM-COV), and 1×Sypro Orange dye was dispensed into the 96-well plate. The same volume of DMSO was used as control. The plates then were heated in a cycler from 25 to 95 °C (71 heating cycles in 35.5 min). The data were analyzed by BIO-RAD software. S6

Cellular Thermal Shift Assay (CETSA). CETSA was performed to detect the ability of compounds interacting with its targets at cellular level. First, the cultured cells were treated with cell media containing 1% DMSO or 10 μ M compounds DI-591, DC-2, and DC-2N for 3–5 h, respectively, the cells were collected, aliquoted into seven PCR tubes, and then heated for 10 min from 43 to 61 °C. As for the dose-dependent assay, 1% DMSO, 0.3, 1, 3, 10 μ M DC-2, or 10 μ M DC-2N was added into the cells for 3–5 h and then heated for 10 min at 55 °C. After that, the cells were lysed by three repeated cycles of freeze—thawing, using liquid nitrogen. Finally, after being centrifuged at 12 000g for 20 min, the supernates were used for Western Blot analysis.

Molecular Docking. The crystal structure (PDB ID: 3TDU, resolution of 1.5 Å) of human Cul1WHB-Dcn1P-acetylated Ubc12N complex was obtained from the RCSB protein data bank. The preparation of protein structure was performed under the Amber 10: EHT force field, using the Quickprep module with the default parameters, which included the addition of hydrogen atoms, the deletion of solvent, the repair of the missing residues, and the setup of the protonation states of the ionizable residues with $pK_a = 7$. The ligand was prepared under Amber 10: EHT force field by energy minimization and conformational search. Afterward, we got the conformational database of the ligand structure for further docking study. All of the conformations then were docked into the binding pocket of human DCN1 with the default parameters, using the method of molecular docking. The docking structure was scored by GBVI/WSA dG and held 20 docking poses in the case of parameters by default.

Coimmunoprecipitation Assay. H1975 cells were treated with 1% DMSO, $10~\mu\mathrm{M}$ compound DC-2N, DC-2, or NAcM-COV, respectively, for 5 h. The cells then were collected and lysed with RIPA buffer. The whole cell lysate was incubated by UBE2M antibody and protein A/G PLUS-Agarose (sc-2003, Santa Cruz Biotech, CA). DCN1 and UBE2M protein associated with beads was eluted by heating and detected by immunoblotting.

Cell-Free NAE Activity Assay. NAE activity assay (in vitro NEDD8~UBE2M formation assay) was conducted according to the manufacturer's instructions. In brief, NEDD8 E1 was mixed with NEDD8, UBE2M, and 10 and 20 μ M MLN4924 or 1, 10, and 100 μ M DC-2 in the reaction buffer and incubated at room temperature for 10 min. The reaction was initiated by the addition of Mg-ATP solution. After 60 min of incubation at 37 °C, the reaction mixtures were added with 1 mM DTT and protein loading buffer (no DTT). The mixtures were then separated by a 12%SDS-PAGE gel. 76

Cell Viability Assay. 3-(4,5-Dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) was used to measure the cell viability, according to the manufacturer's instructions. After the cells were seeded into 96-well plates at a concentration of (2–5) \times 10³ cells per well, serial dilutions of compounds were added and cultured for 72 h. Next, MTT solution (5 mg/mL in PBS) was added and incubated at 37 °C for 4 h. After 150 μ L of DMSO was added followed by removing the MTT, the plate was measured at the absorbance of 490 nm. The data were analyzed using SPSS 20 software.

Colony Formation Assay. First, H1975, PC9, and A549 cells were seeded at a concentration of 1×10^3 cells per well and cultured overnight. The fresh medium then was added with compound **DC-2** into the wells. After 7 days of incubation, the cells were fixed and stained by 0.1% crystal violet solution and then imaged by microscopy (Nikon). Subsequently, the crystal violet crystals were dissolved and measured by a BioTek microplate reader. The data were analyzed using the GraphPad Prism 5 software.

Analysis of Cellular Apoptosis. For the analysis of apoptosis by flow cytometry, first, H1975, PC9, and A549 cells were seeded into 6-well plates ((1–2) \times 10 6 cells/mL) and incubated overnight. Compound DC-2 at 0, 2.5, 5, and 10 $\mu\rm M$ was then added. After treatment for 48 h, the cells were harvested and stained using the Annexin-V-FITC/PI apoptosis kit for 30 min. The cells were then collected and measured by a FACSCalibur flow cytometer (BD Biosciences). All data were analyzed by FlowJo-V10 and GraphPad Prism 5 software.

Western Blot Analysis. H1975 and PC9 cells were seeded in 100 mm² plastic dishes (1 × 106 cells/well) and incubated overnight prior to the addition of compound DC-2 at 0, 0.3, 1, 10, and 20 μ M for 24 h. The cells then were harvested, lysed, and centrifuged at 12 000g for 20 min at 4 °C. The supernatant was determined by a BCA Protein Assay kit, denatured at 100 °C for 10 min, and incubated at -20 °C for Western Blot. After being separated by SDS-PAGE, the protein was transferred to 0.22 μ m nitrocellulose membranes and blocked by 5% skim milk. The membranes then were probed with appropriate primary antibodies and incubated at 4 °C overnight, followed by the treatment of horseradish-peroxidase-conjugated secondary antibody (1:10 000). Subsequently, the membranes were washed with PBST

and examined by enhanced chemiluminescence. The data were determined and analyzed using ImageJ software and GraphPad Prism 5 software.

Water Solubility Assay. Water solubility assay was performed as previous described. 30,73 Briefly, compound DC-2 was added as powder into an Eppendorf tube, which was full of phosphate buffer (PBS, pH 7.4), until a heterogeneous suspension was obtained. The suspension then was sonicated in a water bath for 30 min and shaken for 24 h at room temperature until reaching thermodynamic equilibrium. Suspensions were then centrifuged at 12 000g for 15 min, and the supernatants were filtered, using 0.45 μ m membrane. PBS was used as a control. The filtrate was measured by UV-2700 spectroscopy (Shimadzu, Japan).

siRNA Experiments. Cells were seeded in six-well plates for 16-24 h before siRNA treatment. Medium then was taken away, and the cells were transfected with siRNA using Lipofectamine RNAiMAX Reagent according to the manufacturer's instructions. Three days later, the cells were collected and used for Western Blot assay. Three different siRNA duplexes targeting three specific sequences of DCN1 and a negative control siRNA were synthesized by GenePharma (Shanghai, China). The sense strand nucleotide sequence for DCN1 #1 siRNA was 5'-GGAUAAAGUUCGUCAGUUUTT-3', and the antisense sequence was 5'-AAACUGACGAACUUUAUCCTT-3'; DCN1 #2 siRNA was 5'-GGACAGGAAGAAGUUAGAATT-3' and the antisense sequence was 5'-UUCUAACUUCUUCCU-GUCCTT-3'; DCN1 #3 siRNA was 5'-GCCAUUGCCUACUG-GAACUTT-3', and the antisense sequence was 5'-AGUUCCA-GUAGGCAAUGGCTT-3'. The negative control siRNA (siNeg) sense sequence was 5'-UUCUCCGAACGUGUCACGUTT-3', while the antisense sequence was 5'-ACGUGACACGUUCGGAGAATT-

Statistical Analysis. Data were expressed as mean \pm SD. Statistical differences in two groups were performed by student's *t*-test using GraphPad Prism 5 software. One-way ANOVA analysis was used for multiple group comparison by SPSS 20 software. The *P*-values of 0.05 or less were considered statistically significant.

ASSOCIATED CONTENT

S Supporting Information

The Supporting Information is available free of charge on the ACS Publications website at DOI: 10.1021/acs.jmed-chem.9b00003.

Purities and abundances of all of the recombinant proteins; direct interaction of biotin-AcUBE2M¹⁻²¹ peptides with wild-type DCN1 (GST-DCN1) and its mutants (P97T, F164S, and Y181I); solubility of compound **DC-2** and HPLC spectra of representative compounds; and coordinates of modeled structures of compound **DC-2** in complex with DCN1 after molecular dynamics simulations (PDB code: 3TDU) (PDF) Molecular formula strings (CSV)

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Notes

The authors declare no competing financial interest.

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ABBREVIATIONS USED

DCN1, defective in cullin neddylation 1; UBE2M, NEDD8-conjugating enzyme Ubc12; SCC, squamous cell carcinoma; CRL, cullin ring ligases; NEDD8, neural precursor cell expressed, developmentally down-regulated 8; NAE, NEDD8-activating enzyme; RBX1, ring-box 1; E3, ubiquitin protein ligase; CUL1, cullin 1; CUL2, cullin2; CUL3, cullin3; CUL4, cullin4; CUL5, cullin5; MTT, 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide; CETSA, cellular thermal shift assay; FP, fluorescence polarization; IPTG, isopropyl- β -D-galactopyranoside; HTRF, homogeneous time-resolved fluorescence; NRF2, the nuclear factor-erythroid 2 related factor 2; SAR, structure—activity relationship; NQO1, NADPH: quinone oxidoreductase-1; HO-1, heme oxygenase-1

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