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Research paper

Fluoropyrimidin-2,4-dihydroxy-5-isopropylbenzamides as antitumor agents against CRC and NSCLC cancer cells



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ABSTRACT

A major cause of failure of therapy in patients with non-small cell lung cancer (NSCLC) is development of acquired drug resistance leading to tumor recurrence and disease progression. In addition to the development of new generations of epidermal growth factor receptor-tyrosine kinase inhibitors (EGFR-TKIs), different molecular targets may provide opportunities to improve the therapeutic outcomes. In this study, we utilized the core structure 5-fluorouracil (5-FU) or tegafur, a 5-FU prodrug combined through different linkers with resorcinol to generate a series of fluoropyrimidin-2,4-dihydroxy-5-isopropylbenzamides which inhibit potent Heat Shock Protein 90 (HSP90). These compounds were found to show significant antiproliferative activity in colorectal cancer (CRC) HCT116 and NSCLC A549, H460, and H1975 (EGFR L858R/T790 M double mutation) cells. Compound 12c, developed by molecular docking analysis and enzymatic assays exhibits promising inhibitory activity of HSP90. This compound, 12c shows the most potent HSP90 inhibitory activity with an IC₅₀ value of 27.8 ± 4.4 nM, superior to that of reference compounds AUY-922 (Luminespib) and BIIB021 whose IC_{50} values are 43.0 \pm 0.9 nM and 56.8 \pm 4.0 nM respectively. This strong HSP90 inhibitory activity of **12c** leads to rapid degradation of client proteins EGFR and Akt in NSCLC cells. In addition, **12c** induces significant accumulation of a sub-G1 phase population in parallel with apoptosis by showing activated caspase-3, -8 and -9 and PARP induction. These results provide a new strategy for development of novel HSP90 inhibitors for cancer treatment.

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1. Introduction

Heat shock proteins (HSP) function as molecular chaperones that are responsible for proper folding, assembly, translocation, and degradation of proteins during essential cellular growth and development [1]. HSPs are widely expressed in plants and animals

https://doi.org/10.1016/j.ejmech.2020.112540 0223-5234/© 2020 Elsevier Masson SAS. All rights reserved. and are classified into different families based on their molecular weight [2]. Among HSP families, the role of HSP90 is the most characterized and well-studied. HSP90 is physically associated with numerous co-chaperones such as HSP70, to recruit and interact with diverse substrate or client proteins such as kinases, leading to regulation of cellular processes [3]. In addition, HSP90 has been widely reported to be involved in many human diseases, including neurodegenerative diseases, senescence, cancer and infectious diseases [4]. A number of HSP90 client proteins such as Akt, focal adhesion kinase (FAK) and transcription factor hypoxia-inducible factor 1α (HIF1 α) are involved in tumor proliferation and metastasis [5]. Increased expression of HSP90 is easily detected in many malignancies and is associated with poor prognosis [6]. Consequently, targeting of HSP90 can be effective for cancer treatment

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since HSP90 inhibition will result in interruption of many crucial signaling pathways that are crucial to cancer cell survival. Several HSP90 inhibitors have been developed and are divided into different subgroups based on their structures. For example, resorcinol-based HSP90 inhibitors can be found in a number of clinical trials subjects such as AUY-922 (1. Luminespib), STA-9090 (2. Ganetespib) and AT-13387 (3. Onalespib) (Fig. 1) [7]. These resorcinol-containing compounds have been reported to be active against NSCLC, CRC and other different types of tumors [8].

5-Fluorouracil (4, 5-FU), an analogue of uracil (5) with a fluorine atom at the C-5 position, is converted to several active metabolites to further replace the normal uracil and interrupt the nucleotide synthesis which results in DNA damage in cancer cells [9]. Currently, FOLFIRI a combination of folinic acid (8), 5-FU and irinotecan (6) and FOLFOX (folinic acid, fluorouracil, oxaliplatin (7)) are approved for clinical CRC regimens [10]. Tegafur (9) and capecitabine (10) are prodrugs of 5-FU which are converted intracellularly to active metabolites leading to potent cytotoxicity [11]. Previous reports have suggested that combination therapy of standard regimen FOLFIRI with cetuximab or bevacizumab may improve outcomes in patients with metastatic colorectal cancer [12] and several studies have shown that administration to NSCLC patients of tegafur or capecitabine in combination with other drugs helps to increase patient survival rates [13]. These results suggest one-compound-multi-target agents could be effective and they represent a promising strategy for CRC and NSCLC treatment.

The major clinical challenge in treatment of NSCLC is the development of acquired resistance to drugs [14]. The best study of acquired resistance to epidermal growth factor receptor (EGFR)-TKI is a secondary mutation (T790 M) in exon 20 of EGFR in patients with NSCLC [15]. In order to overcome the resistance to EGFR-TKIs, different types of TKIs and other molecular targeted therapies have been developed with a view to extending disease control [16]. MET amplification, overexpression of hepatocyte growth factor (HGF) and activation of the insulin-like growth factor 1 receptor (IGF1R) have been identified as crucial mechanisms underlying acquired



9, Tegafur

HC Fig. 1. Reported chemotherapeutic agents and HSP90 inhibitors.

resistance to EGFR-TKIs [17]. Previous reports also indicate that EGFR-TKIs, in combination with other targeted drugs such as VEGF, MEK/ERK or HER2 have shown some potential ability to overcome resistance mediated by different mechanisms [18].

Based on previous studies, inhibitors of 5-FU and HSP90 have shown some success in patients with NSCLC [19]. Therefore, we combined the structure of tegafur or 5-FU with resorcinol to exert cytotoxic and HSP90 inhibitory activity and so decrease oncogenic client proteins and enhance the anticancer activity for NSCLC therapy (Fig. 2). A series of compounds (12–13) was synthesized (Fig. 3) and their biological assays are discussed below.

2. Results and discussion

2.1. Chemistry

Tagafur (**6**) reacted with 4-nitrobenzyl bromide in the presence of K_2CO_3 , yielding compound **14** (Scheme 1). Compound **14** with a nitro group was reduced with iron powder and ammonium chloride in isopropanol and water to afford the amino compound (**15**). Compound **15** underwent amide coupling with 1-ethyl-3-(3dimethylaminopropyl)carbodiimide hydrochloride (EDC·HCl), 1hydroxybenzotriazole (HOBt), N-methylmorpholine (NMM) and 2,4-bis(benzyloxy)-5-isopropylbenzoic acid to produce compound **16**. The protected compound (**16**) was debenzylated by Pd/C and hydrogen affording compound **12a**. Compounds **12b-12h** and compounds **13a-13c** were prepared similarly from compound **16** and various substituted groups under basic conditions. The deprotection can be accomplished with hydrogen gas and palladium or with boron trichloride, as is shown in Scheme 2.

2.2. Biological evaluation

2.2.1. HSP90 inhibition

We first evaluated the HSP90 inhibitory activity of synthetic compounds 12a-12h and 13a-13c together with reference compounds AUY-922 and BIIB021, as shown in Table 1. Compound 12a. lacking substitution at the amide linkage, showed no HSP90 inhibitory activity. However, compounds 12b-12h and 13a-13c with the N-substituted groups displayed good inhibitory activity against HSP90. Most of synthetic compounds have IC₅₀ values against HSP90 of ~50 nM which is comparable to those of the reference compounds. Compound **12c** with an ethyl group showed the strongest HSP90 inhibitory activity, suggesting the activity is related to the length of substituent group. The N-substituted benzyl group compounds 12f-12h, and 13a-13c, also displayed significant activity and showed inhibitory activity comparable to that of the reference compounds. Among the synthesized compounds, compound 12c exhibited remarkable HSP90 inhibitory activity with an $IC_{50} = 27.8$ nM, which is about two-fold stronger than the reference compounds.

2.2.2. In vitro cell growth inhibitory activity

In an attempt to evaluate the effect of the synthetic compounds on growth inhibition of cancer cells, we examined synthesized compounds **12a-12h** and **13a-13c** for anti-proliferative activity against CRC HCT116 cells (Table 2). Compared to compound **4** (5-FU), most of synthesized compounds displayed potent inhibitory activities in CRC cells. In addition, the observed pattern of inhibitory activity is consistent with the HSP90 inhibition shown in Table 1. Compound **12c** exhibited anticancer activity with a GI_{50} value of 10 \pm 1 nM which is 15 times more inhibitory than reference



11, R = H or 2-tetrahydrofuran

Fig. 2. Rational design of target compounds.





^aReagents and condition: (a) 4-nitrobenzyl bromide, K₂CO₃, DMF, rt; (b) Iron powder, NH₄Cl, IPA/H₂O, reflux; (c) EDC·HCl, HOBt, NMM, 2,4-bis(benzyloxy)-5-isopropylbenzoic acid , DMF, rt; (d) H₂, 10% Pd/C, MeOH, rt

Scheme 1. Synthetic route of compound 12a^{aa}Reagents and condition: (a) 4-nitrobenzyl bromide, K₂CO₃, DMF, rt; (b) Iron powder, NH₄Cl, IPA/H₂O, reflux; (c) EDC+HCl, HOBt, NMM, 2,4-bis(benzyloxy)-5-isopropylbenzoic acid, DMF, rt; (d) H₂, 10% Pd/C, MeOH, rt.

compound BIIB021.

2.2.3. In vitro cell growth inhibitory activity against human lung cancer cell lines

It has been reported that HSP90 inhibitors such as STA-9090 and AUY-922 display good activity against lung cancer cells [20]. Therefore we also investigated the effect of synthetic compounds against various human lung cancer cell lines, including two carrying wild-type EGFR (A549 and H460), and one EGFR L858R/T790 M double mutations (H1975). As shown in Table 3, a majority of our synthetic compounds have shown the ability to suppress cell proliferation in these three human NSCLC cells independently of their EGFR status. Compounds **13a-13c** exhibit more potent cell growth inhibitory activity than compounds **12b-12h**, suggesting that 5-FU plus resorcinol generates higher cytotoxicity *in vitro* than tegafur-resorcinol. It is surprising that the H1975 cell line is the most sensitive to many of the compounds among **12c-12h**, suggesting that if R₁ is benzyl, cyanobenzyl or chlorobenzyl the

compound may provide some selectivity toward H1975 cells with the EGFR L858R/T790 M double mutation. Compound **12c**, with the best activity against CRC HCT116 cells, as shown in Table 2, also demonstrated promising antiproliferative activity with GI₅₀ values of 0.07, 0.06, and 0.04 μ M against A549, H460 and H1975 NSCLC cells, respectively.

2.2.4. Colony formation in evaluation of antitumor activity in different cancer cells

We examined the colony forming ability of compound **12c** in cells of three different cancers. Although compound **12c** exhibits better antiproliferative activity in CRC HCT116 cells than other tumor cells (Tables 2 and 3), the colony-forming results showed a better inhibitory effect toward NSCLC A549 and H1975 cells with the same concentrations (0.009 μ M and 0.018 μ M) of **12c** (Fig. 4A–C). Taken together, these results show compound **12c** caused a significant inhibition of cell growth and colony-forming ability in NSCLC cells, suggesting **12c** may provide potentially



^aReagents and condition: (a) i. NaH, various substituted group, DMF, rt; ii. H₂, 10% Pd/C, MeOH, rt,
(b) i. NaH, various substituted group, DMF, rt; ii. BCl₃, CH₂Cl₂, 0°C to rt

Scheme 2. Synthetic route of compounds 12b-12h and 13a-13c^a.

^aReagents and condition: (a) i. NaH, various substituted group, DMF, rt; ii. H₂, 10% Pd/C, MeOH, rt, (b) i. NaH, various substituted group, DMF, rt; ii. BCl₃, CH₂Cl₂, 0 °C to rt.

Table 1 HSP90 α inhibition of synthetic compounds (12a-12h and 13a-13c) and reference compounds.

Con	npounds	$\text{HSP90} \alpha \text{ IC}_{50} (n\text{M} \pm \text{SD}^{\text{a}})$	Compounds	HSP90 α IC ₅₀ (nM ± SD ^a)
12a		>1000	12h	58.8 ± 6.9
12b 12c		40.9 ± 19.3 27.8 + 4.4	13a 13b	44.9 ± 0.8 48.0 ± 1.5
12d		48.2 ± 13.8	13c	54.4 ± 0.6
12e 12f		45.3 ± 5.0 45.3 ± 7.0	AUY-922 (1) BIIB021	43.0 ± 0.9 56.8 + 4.0
12g		54.7 ± 16.3		

^a SD: standard deviation. All experiments were independently performed at least three times.

useful antitumor activity for NSCLC treatment.

2.2.5. Evaluation of expression level of Hsp90 client proteins

Next, we examined some critical biomarkers of Hsp90 inhibition in A549 and H1975 cells. Inhibition of HSP90 will lead to heat shock factor 1 (HSF1) activation, consequently inducing several small heat shock proteins to help appropriate folding of misfolded proteins [21]. As shown in Fig. 5A and B, treatment with compound **12c** greatly induced the expression level of HSP70 without affecting HSP90 levels. Furthermore, several client proteins such as protein kinase B (akt) and EGFR were also downregulated in response to compound **12c**, indicating compound **12c** achieves significant suppression of HSP90 activity in cells.

2.2.6. Evaluation of cell cycle progression and cell death response

We examined the effects of compound **12c** on cell cycle progression using flow cytometry. Notably, the subG1 phase showed a dramatic induction after 48 h of treatment (Fig. 6A and B) in parallel with significant activation of caspase-3, -8, and -9, and poly-(ADPribose) polymerase (PARP) (Fig. 7A and B). These results suggest that compound **12c** induces significant apoptotic cell death in NSCLC.

2.2.7. Molecular docking study

Molecular docking analysis was performed to illuminate the interactions between **12c** and HSP90 (PDB ID: 5GGZ). The docking pose of **12c** in HSP90, can be separated into four distinct sites (Fig. 8). In Site 1 (S1), the resorcinol group of **12c** forms hydrogen bonds with residues Leu48, Asp93, and Thr184, and hydrophobic interactions occur between its isopropyl moiety and residues Leu48 and Phe138, while the benzene ring creates hydrophobic interactions with residues Ala55 and Val186. Site 2 (S2) contains an

Table 2

Antiproliferative activity (GI₅₀) of synthetic compounds (12a-12h and 13a-13c) against colorectal HCT116 cancer cell line.

Compounds	HCT116 GI ₅₀ (μ M ± SD ^a)	Compounds	HCT116 GI ₅₀ (μ M \pm SD ^a)
12a	>10	13a	0.030 ± 0.001
12b	0.120 ± 0.010	13b	0.080 ± 0.001
12c	0.010 ± 0.001	13c	0.040 ± 0.006
12d	0.960 ± 0.210	STA-9090 (2)	0.150 ± 0.020
12e	0.780 ± 0.002	5-FU (4)	>10
12f	0.410 ± 0.050	Tegafur (9)	>10
12g	0.700 ± 0.110	BIIB021	0.150 ± 0.020
12h	0.720 ± 0.011		

^a SD: standard deviation. All experiments were independently performed at least three times.

Tabla 2

12e

12f

12g

12h

13a

13b

13c AUY922 (1)

5-FU (4) Tegafur (9)

BIIB021

Antiproliferative activity (GI ₅₀) of compounds (12a-12h and 13a-13c) against lung A549, H460 and H1975 cancer cell line.						
Compounds	A549 GI_{50} ($\mu M \pm SD^{a}$)	H460 GI ₅₀ (μ M ± SD ^a)				
12a	>10	>10				
12b	0.09 ± 0.01	0.07 ± 0.03				
12c	0.07 ± 0.01	0.06 ± 0.01				
12d	0.65 ± 0.03	0.52 ± 0.12				

 0.21 ± 0.10 ^a SD: standard deviation. All experiments were independently performed at least three times.

 0.80 ± 0.07

 0.35 ± 0.04

 0.56 ± 0.03

 0.92 ± 0.11

 0.04 ± 0.00

 0.10 ± 0.01

 0.03 ± 0.00

 0.01 ± 0.00

 19.76 ± 2.9

0.26 + 0.04

>50

N-ethyl moiety that occupies a hydrophobic pocket formed by residues Ala55, Met98 and Tyr61. The phenyl group at Site 3 (S3) occupies a cavity within the binding site and serves as a linker between the ethyl and the tegafur moiety at Site 4 (S4). The tegafur moiety forms hydrogen bonds with residues Asn51 and Phe138. When comparing 12c with its derivatives, 12c has an increased HSP90 inhibitory effect over that of its derivatives, such as unsubstituted **12a** and **12b** with a methyl group in place of the ethyl moiety. These differences reduce their inhibitory effect against HSP90 (Table 1). Consequently, it could be concluded that the ethyl moiety in 12c is important for its activity against HSP90.

To further elucidate interactions between compound 12c and HSP90, we compared the compound to the co-crystal ligand, 6 TN. We observed an overlap at Site 1 and Site 2 (Fig. 8C). This co-crystal ligand also contains a resorcinol group at Site 1. Hydrogen bonds form with residues Asp93 and Thr184 at Site 1. Residues Thr184 and Ala55 form hydrophobic interactions with the benzene ring. An isopropyl moiety is attached to the benzene ring and forms further hydrophobic interactions with residues Phe138 and Val150. The cocrystal ligand also contains a pyridopyrimidine structure. A part of this moiety occupies Site 2, forming hydrophobic interactions with residues Ala55, Met98 and Leu107. Because of its position, the cocrystal ligand does not align with Site 3 and Site 4. The reported IC₅₀ value of co-crystal ligand is 56 nM [22]. When compared to compound 12c, the difference in potency may be due to its interactions at Site 3 and Site 4.

We previously found that compound **12c** has a comparable IC₅₀ value to the HSP90 inhibitor AUY-922 (Table 1). A co-crystal structure of HSP90 with AUY-922 has been determined previously (PDB ID: 2VCI) [23]. As a result, we compared compound 12c to the AUY-922 co-crystal ligand. We found that both compounds align with the four distinct sites (Fig. 8D). AUY-922 also contains a resorcinol and isopropyl moiety that occupies Site 1, similar to compound **12c**. Hydrogen bonds are formed with residues Asn51, Asp93 and Thr184. The isopropyl moiety occupies a similar hydrophobic pocket by residues Leu107 and Phe138. At Site 2, residues Lys58 and Gly97 form hydrogen bonds to the oxygen and nitrogen of the N-ethylformamide, respectively. Further hydrophobic interactions by residue Ala55 occur at Site 3. However, Site 4 did not show stable interactions between AUY-922 and HSP90. The additional hydrogen bond at Site 3 and hydrophobic interactions at Site 4 suggests areas that can be exploited for HSP90 inhibition. Together, the molecular docking analysis is consistent with the potency of 12c against HSP90.

3. Conclusion

 0.40 ± 0.19

 0.20 ± 0.08

 0.46 ± 0.08

0.39 + 0.16

 0.03 ± 0.01

 0.07 ± 0.01

 0.07 ± 0.02

 0.03 ± 0.01

In this study, a series of fluoropyrimidin-2,4-dihydroxy-5isopropylbenzamides (12–13) were synthesized as HSP90 inhibitors in an attempt to explore the compounds' biological activity against CRC and NSCLC. Among all the synthesized compounds, compound 12c displays remarkable activity with an IC₅₀ value of 27.8 nM in inhibition of HSP90 activity. In in vitro cell growth inhibition, 12c not only shows good activity against HCT116 cancer cell with a GI₅₀ value of 0.01 μ M but also exhibits good activity with GI₅₀ values of 0.07, 0.06, and 0.04 µM against A549, H460 and H1975 human NSCLC cell lines, respectively. In addition, compound 12c significantly reduces the client proteins EGFR and Akt and induces apoptotic cell death in NSCLC cells. In this study, we have developed **12c** as a potential agent and a new strategy for development of novel HSP90 inhibitors for the treatment of cancer.

H1975 GI₅₀ ($\mu M \pm SD^{a}$)

>10 0.06 ± 0.00 0.04 ± 0.00 0.34 ± 0.03

 0.16 ± 0.04

 0.13 ± 0.01

 0.16 ± 0.03

0.23 + 0.05

0.03 + 0.00

 0.05 ± 0.02

 0.02 ± 0.00

 0.02 ± 0.00

0.20 + 0.06

4. Experimental section

4.1. Chemistry

Nuclear magnetic resonance (¹H and ¹³C NMR) spectra were obtained with Bruker DRX-500 spectrometer operating at 500 and 125 MHz and Bruker Fourier 300 and 75 MHz. Chemical shifts are reported in parts per million (ppm, δ) downfield from TMS as an internal standard. High-resolution mass spectra (HRMS) were measured with AB SCIE X (QSTAR® XL) High Resolution Electrospray (ESI) Mass Spectrometry spectrometer. Melting points were measured with Buchi B-545 (Buchi, Switzerland). Purity of the final compounds was achieved with a Waters Acquity UPLC system using C-18 column (Waters Acquity UPLC BEH C18, 1.7 µm, 2.1 mm \times 50 mm). Flash column chromatography used silica gel: SILICYCLE (SiliaFlash Irregular Silica Gel P60, 40–63 µm, 60 Å (R12030B).

4.1.1. N-(4-((5-Fluoro-2,6-dioxo-3-(tetrahydrofuran-2-yl)-3,6dihydropyrimidin-1(2H)-yl)methyl)phenyl)-2,4-dihydroxy-5isopropylbenzamide (12a)

A mixture of 16 (2.05 mmol), 10% palladium on carbon (0.4 g) in MeOH (40 ml) was stirred at room temperature (rt) under hydrogen overnight. The organic layer was filtered and the residue was purified by flash chromatography over silica gel to afford compound **12a** in 46% yield. ¹H NMR (500Hz, DMSO- d_6) δ (ppm): 1.17 (d, *J* = 6.9Hz, 6H), 1.90–1.93 (m, 2H), 2.02–2.04 (m, 1H), 2.21–2.28 (m, 1H), 3.07–3.13 (m, 1H), 3.81 (q, J = 7.5Hz, 1H), 4.22–4.26 (m, 1H),



Fig. 4. Compound **12c** suppresses colony formation in colorectal (CRC) and non-small cell lung cancer (NSCLC) cells. (A–C) Compound **12c**-mediated concentration-dependent inhibition of colony formation in three different tumor cells lines (left panel) and quantification of colony numbers, expressed as a percentage of controls (right panel). Data are expressed as means \pm S.D. of at least three independent experiments. *, *P* < 0.05; **, *P* < 0.01; and ***, *P* < 0.001 compared with the control group.

4.95 (d, J = 4.5Hz, 2H), 5.95–5.97 (m, 1H), 6.35 (s, 1H), 7.29 (d, J = 8.5Hz, 2H), 7.55 (d, J = 8.5Hz, 2H), 7.83 (s, 1H), 7.97 (d, J = 7Hz, 1H), 10.09 (s, 1H), 10.16 (s, 1H), 12.03 (s, 1H). ¹³C NMR (125 MHz, DMSO- d_6) δ (ppm): 22.64, 23.47, 26.15, 31.65, 43.69, 69.56, 87.47, 102.57, 107.37, 121.34, 123.83, 124.10, 126.43, 128.25, 132.00, 137.43,

138.51, 140.33, 148.84, 156.51, 156.71, 159.13, 159.70, 167.22. $mp=222.6-223.2\ ^{\circ}C.$ HRMS (ESI) for $C_{25}H_{27}FN_{3}O_{6}\ [M+H]^{+}:$ Calcd, 484.1878; Found, 484.1884.



Fig. 5. Compound 12c reduces HSP90 client proteins. (A–B) Compound 12c suppresses protein expression level of HSP90 client proteins in NSCLC A549 (A) and H1975 (B) cells. Cells were treated with indicated concentrations for 24 h, and cell lysates were immunoblotted using the indicated antibodies.



Fig. 6. Compound **12c** induces subG1 phase accumulation in NSCLC cells. (A–B) Concentration-dependent effects of Compound **12c** on subG1 phase accumulation in A549 (A) and H1975 (B) cells. Cells were treated with DMSO or Compound **12c** with indicated concentrations for the indicated times, and the cell cycle distribution was analyzed by flow cytometry. The right panel are quantitative data based on flow cytometry histograms, and are presented as means ± S.D. of at least three independent experiments that yielded similar results.



Fig. 7. Compound **12c** induces apoptosis in NSCLC. (A–B) Effects of compound **12c** on apoptosis in NSCLC cells. compound **12c** increased levels of the cleaved (activated) forms of PARP, γH2AX, caspase-3, -8, and -9 in a concentration-dependent manner. Cells were exposed to compound **12c** with indicated concentrations for 48 h, and cell lysates were immunoblotted using the indicated antibodies.

4.1.2. N-(4-((5-Fluoro-2,6-dioxo-3-(tetrahydrofuran-2-yl)-3,6-dihydropyrimidin-1(2H)-yl)methyl)phenyl)-2,4-dihydroxy-5-isopropyl-N-methylbenzamide (**12b**)

A mixture of 16 (2.10mmole), NaH (2.52mmole), MeI (2.52mmole) and DMF (3 ml) was stirred at rt for 2 h. The reaction was guenched with water and extracted by EtOAc. The residue was purified by flash chromatography over silica gel to afford an intermediate. Then a mixture of this intermediate, 10% palladium on carbon (0.4 g) in MeOH (40 ml) was stirred at rt under hydrogen overnight. The organic layer was filtered and the residue was purified by flash chromatography over silica gel to afford compound **12b** in 42% yield. ¹H NMR (500 MHz, DMSO- d_6) δ (ppm): 0.63 (d, J = 3Hz, 3H), 0.65 (d, J = 3Hz, 3H), 1.89–1.96 (m, 2H), 1.97–2.01 (m, 1H), 2.20-2.26 (m, 1H), 2.75-2.80 (m, 1H), 3.29 (s, 3H), 3.80 (q, J = 7.5Hz, 1H), 4.24 (q, J = 6Hz, 1H), 4.91 (q, J = 14.5Hz, 2H), 5.92–5.94 (m, 1H), 6.18 (s, 1H), 6.45 (s, 1H), 7.13 (d, J = 8.5Hz, 2H), 7.29 (d, J = 8Hz, 2H), 7.95 (d, J = 6.5Hz, 2H), 10.72 (s, 1H). ¹³C NMR (125 MHz, DMSO- d_6) δ (ppm): 22.07, 23.45, 24.96, 31.67, 38.17, 43.49, 69.59, 87.45, 102.27, 109.06, 123.76, 124.03, 124.63, 126.50, 127.72, 129.22, 134.58, 138.54, 140.35, 144.56, 148.77, 156.37, 156.57, 157.55, 157.85, 170.29. mp = 199.3-199.9 °C. HRMS (ESI) for C₂₆H₂₉FN₃O₆ [M+H]⁺: Calcd, 498.2035; Found, 498.2041.

4.1.3. N-Ethyl-N-(4-((5-fluoro-2,6-dioxo-3-(tetrahydrofuran-2-yl)-3,6-dihydropyrimidin-1(2H)-yl)methyl)phenyl)-2,4-dihydroxy-5-isopropylbenzamide (12c)

The title compound was obtained as a solid in 33% yield from compound **16** in a manner similar to that described for the preparation of **12b.** ¹H NMR (500 MHz, DMSO- d_6) δ (ppm): 0.60 (d,

J = 3.1Hz, 3H), 0.61 (d, *J* = 3Hz, 3H), 1.06 (t, *J* = 7Hz, 3H), 1.89–1.92 (m, 2H), 1.97–2.00 (m, 1H), 2.20–2.28 (m, 1H), 2.72–2.78 (m, 1H), 3.76–3.83 (m, 3H), 4.23–4.27 (m, 1H), 4.92 (q, *J* = 14.5Hz, 2H), 5.92–5.94 (m, 1H), 6.17 (s, 1H), 6.42 (s, 1H), 7.12 (d, *J* = 8Hz, 2H), 7.32 (d, *J* = 8.5Hz, 2H), 7.95 (d, *J* = 6.5Hz, 2H), 9.76 (s, 1H), 10.93 (s, 1H). ¹³C NMR (125 MHz, DMSO-*d*₆) δ (ppm): 12.52, 22.02, 23.44, 24.91, 31.67, 43.49, 45.13, 69.58, 87.44, 102.25, 123.75, 124.01, 124.54, 127.43, 127.75, 129.36, 134.81, 138.54, 140.36, 142.81, 148.77, 156.35, 156.56, 157.63, 158.44, 169.98. mp = 173.4–174.6 °C. HRMS (ESI) for C₂₇H₃₁FN₃O₆ [M+H]⁺: Calcd, 512.2191; Found, 512.2195.

4.1.4. N-(4-((5-Fluoro-2,6-dioxo-3-(tetrahydrofuran-2-yl)-3,6dihydropyrimidin-1(2H)-yl)methyl)phenyl)-2,4-dihydroxy-5isopropyl-N-propylbenzamide (**12d**)

The title compound was obtained as a solid in 42% yield from compound **16** in a manner similar to that described for the preparation of **12b.** ¹H NMR (500 MHz, DMSO-*d*₆) δ (ppm): 0.60 (d, J = 3.4Hz, 3H), 0.62 (d, J = 3.4Hz, 3H), 0.82 (t, J = 7Hz, 3H), 1.46–1.50 (m, 2H), 1.89–1.92 (m, 2H), 1.99–2.00 (m, 1H), 2.22–2.26 (m, 1H), 2.49–2.77 (m, 1H), 3.71 (t, J = 7.5H, 2H), 3.80 (q, J = 7.5Hz, 1H), 4.23–4.25 (m, 1H), 4.91 (q, J = 14.5Hz, 1H), 5.92–5.94 (m, 1H), 6.17 (s, 1H), 6.41 (s, 1H), 7.12 (d, J = 8Hz, 2H), 7.30 (d, J = 8.5Hz, 2H), 7.95 (d, J = 6.5Hz, 2H), 10.82 (s, 1H). ¹³C NMR (125 MHz, DMSO-*d*₆) δ (ppm): 11.13, 14.08, 20.29, 22.05, 23.46, 24.93, 31.69, 43.50, 51.50, 69.60, 87.46, 102.27, 108.94, 123.76, 124.03, 124.56, 127.30, 127.66, 129.30, 134.73, 138.55, 140.37, 143.00, 148.79, 156.37, 156.58, 157.56, 158.20, 170.11 mp = 102.6–103.5 °C. HRMS (ESI) for C₂₈H₃₃FN₃O₆ [M+H]⁺: Calcd, 526.2348; Found, 526.2354.



Fig. 8. Docking pose of compound **12c** in HSP90. (A) **12c** (blue) occupies the HSP90 (gray, PDB ID: 5GGZ) binding site. (B) A 2D representation of **12c** docked in HSP90. Hydrogen bonds are denoted by dashed lines, green lines represent areas of hydrophobic interactions. Interacting residues are labeled as shown. (C) The docking pose of **12c** superimposed with the co-crystal ligand, 6 TN. Overlapping sections are highlighted as previously described. (D) The docking pose of **12c** superimposed with the co-crystal ligand AUY-922 (purple, PDB ID: 2VCI). Interacting residues are labeled and represented as sticks. Hydrogen bonds are denoted by dotted green lines. The four distinct sites of **12c** are colored as purple, red, yellow and blue, which are located at sections **S1**, **S2**, **S3** and **S4**, respectively. (For interpretation of the references to color in this figure legend, the reader is referred to the Web version of this article.)

4.1.5. N-Benzyl-N-(4-((5-fluoro-2,6-dioxo-3-(tetrahydrofuran-2-yl)-3,6-dihydropyrimidin-1(2H)-yl)methyl)phenyl)-2,4-dihydroxy-5-isopropylbenzamide (**12e**)

The title compound was obtained as a solid in 32% yield from compound **16** in a manner similar to that described for the preparation of **12b.** ¹H NMR (500 MHz, DMSO-*d*₆) δ (ppm): 0.64 (d, J = 3Hz, 3H), 0.65 (d, J = 3.5Hz, 3H), 1.88–1.92 (m, 2H), 1.95–1.98 (m, 1H), 2.18–2.24 (m, 1H), 2.75–2.80 (m, 1H), 3.77–3.82 (m, 1H), 4.21–4.25 (m, 1H), 4.85 (q, J = 14Hz, 2H), 5.03 (s, 1H), 5.90–5.92 (m, 1H), 6.20 (s, 1H), 6.51 (s, 1H), 7.05 (d, J = 8.5Hz, 2H), 7.20–7.21 (m, 3H), 7.26–7.29 (m, 4H), 7.93 (d, J = 6.5Hz, 1H), 9.74 (s, 1H), 10.61 (s, 1H). ¹³C NMR (125 MHz, DMSO-*d*₆) δ (ppm): 22.55, 23.91, 25.48, 32.15, 43.93, 53.44, 60.22, 70.05, 87.92, 102.78, 124.21, 124.48, 125.21, 127.45, 127.47, 128.10, 128.79, 129.44, 134.97, 137.93, 139.00, 140.81, 143.36, 149.22, 156.82, 157.02, 158.09, 158.18, 170.84. mp = 99.8–100.7 °C. HRMS (ESI) for C₃₂H₃₃FN₃O₆ [M+H]⁺: Calcd, 574.2348; Found, 574.2353.

4.1.6. N-(4-Cyanobenzyl)-N-(4-((5-fluoro-2,6-dioxo-3-(tetrahydrofuran-2-yl)-3,6-dihydropyrimidin-1(2H)-yl)methyl) phenyl)-2,4-dihydroxy-5-isopropylbenzamide **(12f)**

The title compound was obtained as a solid in 35% yield from compound **16** in a manner similar to that described for the preparation of **12b.** ¹H NMR (500 MHz, DMSO-*d*₆) δ (ppm): 0.67 (d, J = 3Hz, 3H), 0.68 (d, J = 3Hz, 3H), 1.88–1.92 (m, 2H), 1.97–1.98 (m, 2H), 2.18–2.24 (m, 1H), 2.76–2.80 (m, 1H), 3.77–3.82 (m, 1H), 4.21–4.25 (m, 1H), 4.86 (q, J = 14.5Hz, 2H), 5.11 (s, 1H), 5.90–5.91 (m, 1H), 6.19 (s, 1H), 6.55 (s, 1H), 7.09 (d, J = 8.5Hz, 2H), 7.20 (d, J = 8.5Hz, 2H), 7.49 (d, J = 8Hz, 2H), 7.75 (d, J = 8.5Hz, 2H), 7.93 (d, J = 7Hz, 1H), 9.74 (s, 1H), 10.45 (s, 1H). ¹³C NMR (125 MHz, DMSO-*d*₆) δ (ppm): 22.09, 23.42, 25.04, 31.65, 43.43, 52.79, 69.57, 87.43, 102.28, 109.81, 118.75, 123.74, 124.01, 124.83, 126.85, 127.56, 128.43, 128.92, 132.29, 134.61, 138.50, 140.32, 142.66, 143.47, 148.74, 156.34, 156.54, 157.27, 157.59, 170.38. mp = 117.5–118.4 °C. HRMS (ESI) for C₃₃H₃₂FN₄O₆ [M+H]⁺: Calcd, 599.2300; Found, 599.2307.

4.1.7. N-(4-Chlorobenzyl)-N-(4-((5-fluoro-2,6-dioxo-3-(tetrahydrofuran-2-yl)-3,6-dihydropyrimidin-1(2H)-yl)methyl) phenyl)-2,4-dihydroxy-5-isopropylbenzamide **(12g)**

The title compound was obtained as a solid in 34% yield from compound **16** in a manner similar to that described for the preparation of **12b** ¹H NMR (500 MHz, DMSO- d_6) δ (ppm): 0.64 (t, J = 2.1Hz, 6H), 1.89–1.94 (m, 2H), 1.95–1.97 (m, 1H), 2.20–2.24 (m, 1H), 3.79 (q, J = 7.5Hz, 1H), 4.83 (d, J = 14.5Hz, 1H), 4.88 (d, J = 14.5Hz, 1H), 5.01–5.03 (m, 2H), 5.89–5.92 (m, 1H), 6.20 (s, 1H), 6.50 (s, 1H), 7.05 (d, J = 8Hz, 2H), 7.19–7.21 (m, 3H), 7.26–7.33 (m, 4H), 7.92 (d, J = 6.5Hz, 1H). ¹³C NMR (125 MHz, DMSO- d_6) δ (ppm): 22.20, 23.48, 25.06, 31.74, 43.52, 53.05, 69.65, 87.51, 102.37, 123.63, 124.80, 127.05, 127.08, 127.68, 128.38, 129.03, 129.69, 131.39, 134.33, 136.33, 137.15, 138.56, 140.13, 143.21, 148.81, 156.11, 157.84, 170.44. mp = 99.8–101.1C. HRMS (ESI) for C₃₂H₃₂ClFN₃O₆ [M+H]⁺: Calcd, 608.1958; Found, 608.1962.

4.1.8. N-(4-((5-Fluoro-2,6-dioxo-3-(tetrahydrofuran-2-yl)-3,6dihydropyrimidin-1(2H)-yl)methyl)phenyl)-2,4-dihydroxy-5isopropyl-N-(4-methoxybenzyl)benzamide (**12h**)

The title compound was obtained as a solid in 35% yield from compound **16** in a manner similar to that described for the preparation of **12b**. ¹H NMR (500 MHz, DMSO- d_6) δ (ppm): 0.61 (d, J = 3.3Hz, 3H), 0.63 (d, J = 3.3Hz, 3H), 1.89–1.94 (m, 3H), 2,20–2.24 (m, 1H), 2.74–2.77 (m, 1H), 3.77 (s, 3H), 4.21–4.25 (m, 1H), 4.86 (q, J = 14.5, 2H), 4.95 (s, 2H), 5.90–5.91 (m, 1H), 6.19 (s, 1H), 6.47 (s, 1H), 6.81 (d, J = 8.5Hz, 2H), 7.01 (d, J = 8Hz, 2H), 7.16 (d, J = 8.5Hz, 2H), 7.20 (d, J = 8.5Hz, 2H), 7.92 (d, J = 6.2Hz, 2H), 9.78 (s, 1H), 10.70 (s, 1H). ¹³C NMR (125 MHz, DMSO- d_6) δ (ppm): 22.09, 23.4, 25.02, 31.72, 43.52, 55.00, 59.81, 69.64, 87.50, 102.34, 113.74, 123.77, 124.04, 124.75, 127.15, 127.72, 128.31, 129.06, 129.21, 129.29, 134.55, 138.56, 142.84, 148.80, 157.63, 158.35, 170.37. mp = 102.1–102.8 °C. HRMS (ESI) for C₃₃H₃₅FN₃O₇ [M+H]+: Calcd, 604.2454; Found, 604.2466.

4.1.9. *N-Ethyl-N-(4-((5-fluoro-2,6-dioxo-3,6-dihydropyrimidin-1(2H)-yl)methyl)phenyl)-2,4-dihydroxy-5-isopropylbenzamide* (**13a**)

A mixture of 16 (1.85 mmol), NaH (2.22 mmol), MeI (2.22 mmol) and DMF (3 ml) was stirred at 0 °C for 10 min and left at rt for 2 h. The reaction was quenched with water and extracted with EtOAc. The residue was purified by flash chromatography over silica gel to afford an intermediate compound. Then a mixture of the intermediate compound and boron trichloride (1 M) (17% in hexane) in CH₂Cl₂ was stirred at 0 °C for 3 h. The reaction was quenched with water and extracted with CH₂Cl₂. The organic layer was collected and dried over anhydrous MgSO4 and concentrated in vacuo to yield an oily residue which was purified by flash chromatography over silica gel to afford compound **13a** in 36% yield. ¹H NMR (500 MHz, DMSO- d_6) δ (ppm): 0.34 (d, I = 7Hz, 6H), 0.78 (t, I = 6.5 Hz, 3H), 3.51 (q, J = 7Hz, 2H), 4.62 (s, 2H), 5.92 (s, 1H), 6.16 (s, 1H), 6.84 (d, J = 8.5Hz, 2H), 7.04 (d, J = 8Hz, 2H), 7.58 (t, J = 5.5Hz, 1H), 9.53 (s, 1H), 10.70 (s, 1H), 10.91 (d, J = 6Hz, 1H). ¹³C NMR (125 MHz, DMSO- d_6) δ (ppm): 12.94, 22.52, 25.41, 43.35, 45.65, 102.73, 109.05, 125.06, 125.49, 125.74, 127.89, 128.28, 129.79, 135.56, 139.04, 140.84, 143.23, 150.28, 157.59, 157.80, 158.13, 159.01, 170.81. $mp = 114.6 - 115.4 \circ C. HRMS (ESI) for C_{23}H_{25}FN_3O_5 [M+H]^+: Calcd,$ 442.1773; Found, 442.1780.

4.1.10. N-Allyl-N-(4-((5-fluoro-2,6-dioxo-3,6-dihydropyrimidin-1(2H)-yl)methyl)phenyl)-2,4-dihydroxy-5-isopropylbenzamide (13b)

The title compound was obtained as a solid in 42% yield from compound **16** in a manner similar to that described for the preparation of **13a**. ¹H NMR (300 MHz, DMSO- d_6) δ (ppm): 0.35 (d,

J = 6.6Hz, 6H), 3.70 (q, J = 7.2Hz, 2H), 4.08 (d, J = 5.1Hz, 2H), 4.58 (s, 2H), 4.76-4.85 (m, 2H), 5.49-5.62 (m, 1H), 5.89 (s, 1H), 6.18 (s, 1H), 6.79 (d, J = 8.1Hz, 2H), 6.96 (d, J = 8.1Hz, 2H), 7.54 (t, J = 5.7Hz, 1H), 9.47 (s, 1H), 10.41 (s, 1H). ¹³C NMR (125 MHz, DMSO-*d* $_6) <math>\delta$ (ppm): 22.56, 25.46, 43.35, 53.09, 60.23, 102.74, 109.49, 117.84, 125.18, 125.52, 125.77, 127.54, 128.17, 129.48, 133.97, 135.34, 139.02, 140.83, 143.44, 150.28, 157.60, 157.80, 158.09, 158.42, 170.51. mp = 210.3-211.2 °C. HRMS (ESI) for C₂₄H₂₅FN₃O₅ [M+H]⁺: Calcd, 454.1773; Found, 454.1778.

4.1.11. N-(4-((5-Fluoro-2,6-dioxo-3,6-dihydropyrimidin-1(2H)-yl) methyl)phenyl)-2,4-dihydroxy-5-isopropyl-N-(prop-2-yn-1-yl) benzamide (**13c**)

The title compound was obtained as a solid in 47% yield from compound **16** in a manner similar to that described for the preparation of **13a**. ¹H NMR (300 MHz, DMSO-*d*₆) δ (ppm): 0.67 (d, J = 3.9 Hz, 6H), 2.62–2.81 (m, 1H), 3.12 (s, 1H), 4.55 (d, J = 1.2Hz, 2H), 4.89 (s, 2H), 6.19 (s, 1H), 6.49 (s, 1H), 7.14 (d, J = 4.8Hz, 2H), 7.28 (d, J = 5.1Hz 2H), 7.84 (d, J = 3.3Hz 1H), 10.52 (s, 1H). ¹³C NMR (125 MHz, DMSO-*d*₆) δ (ppm): 22.60, 25.48, 43.39, 75.05, 80.15, 102.78, 109.32, 125.60, 125.86, 127.71, 128.14, 129.42, 135.86, 139.02, 140.82, 142.71, 150.30, 157.61, 157.81, 158.21, 158.25, 170.35 mp = 214.2–214.9 °C. HRMS (ESI) for C₂₄H₂₃FN₃O₅ [M+H]⁺: Calcd, 452.1616; Found, 452.1622.

4.1.12. 5-Fluoro-3-(4-nitrobenzyl)-1-(tetrahydrofuran-2-yl) pyrimidine-2,4(1H,3H)-dione (14)

A solution of 5-fluoro-1-(tetrahydrofuran-2-yl)pyrimidine-2,4(1H,3H)-dione (5.00 mmol), 1-(bromomethyl)-4-nitrobenzene (6.50 mmol), K₂CO₃ (6.50 mmol) in DMF (10 ml) was stirred at rt overnight. The reaction was quenched with water and extracted with EtOAc. The residue was purified by flash chromatography over silica gel to afford compound **15** in 70% yield. ¹H NMR (300 MHz, CDCl₃) δ (ppm): 1.95 (m, 1H), 2.06 (m, 2H), 2.40 (m, 1H), 3.97 (q, *J* = 2.1 Hz, 1H), 4.22 (m, 1H), 5.18 (q, *J* = 7.2 Hz, 2H), 5.97 (m, 1H), 7.42 (d, *J* = 5.7 Hz, 1H), 7.62 (d, *J* = 8.7Hz, 2H), 8.16 (d, *J* = 8.7 Hz, 2H).

4.1.13. 3-(4-Aminobenzyl)-5-fluoro-1-(tetrahydrofuran-2-yl) pyrimidine-2,4(1H,3H)-dione (15)

A mixture of **14** (0.69 mmol), iron powder (2.07 mmol), ammonium chloride (1.38 mmol), water (1.4 ml) and isopropyl alcohol (5.6 ml) was stirred reflux for 3 h. The reaction was quenched with water and extracted with EtOAc. The residue was purified by flash chromatography over silica gel to afford **15** in 81% yield. ¹H NMR (300 MHz CDCl₃) δ (ppm): 1.98 (m, 1H), 2.02 (m, 2H), 2.38 (m, 1H), 3.96, (q, *J* = 6.6 Hz, 1H), 4.20 (m, 1H), 5.03 (q, *J* = 4.2 Hz, 2H), 5.95 (m, 1H), 6.94 (d, *J* = 8.4 Hz, 2H), 7.35 (d, *J* = 5.7 Hz, 1H), 7.41 (d, *J* = 8.4 Hz, 2H).

4.1.14. 2,4-bis(Benzyloxy)-N-(4-((5-fluoro-2,6-dioxo-3-(tetrahydrofuran-2-yl)-3,6-dihydropyrimidin-1(2H)-yl)methyl) phenyl)-5-isopropylbenzamide (**16**)

A mixture of 2,4-bis(benzyloxy)-5-isopropylbenzoic acid (0.90 mmol), 1-ethyl-3-(3-dimethylaminopropyl)carbodiimide (1.13 mmol), hydroxybenzotriazole (0.90 mmol), N-methylmorpholine (1.13 mmol) and **15** (0.75 mmol) was stirred at rt overnight. The reaction was quenched with water and extracted with EtOAc. The residue was purified by flash chromatography over silica gel to afford compound **16** in 90% yield. ¹H NMR (300 MHz, CDCl₃): 1.25 (m, 6H), 1.93 (m, 1H), 2.02 (m, 1H), 2.05 (m, 1H), 2.38 (m, 1H), 3.34 (t, J = 6.9 Hz, 1H), 3.95 (q, J = 3.0Hz, 1H), 4.18 (m, 1H), 5.09–5.29 (m, 6H), 5,96 (m, 1H), 6.61 (m, 1H), 7.16–7.63 (m, 15H), 8.18 (s, 1H), 9.85 (s, 1H).

4.2. Biological activity

4.2.1. Cell lines and reagents

A549, H1975, and HCT116 cells were obtained from the American Type Culture Collection (ATCC) (Manassas, VA, USA). Cells were maintained in 10% fetal bovine serum (FBS)-supplemented RPMI 1640 medium (GIBCO, Grand Island, NY, USA) and 1% penicillin–streptomycin (GIBCO) at 37 °C in a humidified incubator containing 5% CO₂. Antibodies against various proteins were obtained from the following sources: PARP (Poly-ADP-ribose polymerase) was obtained from Santa Cruz Biotechnology Inc. (Dallas, TX, USA). Caspase 8, caspase 9, and γ H2AX, were obtained from Cell signaling (Danvers, MA, USA). β -actin and GAPDH were obtained from Millipore (Billerica, MA, USA). Caspase 3 was obtained from Novous (Littleton, CO, USA). Anti-mouse and anti-rabbit IgGs were from Jackson ImmunoResearch Laboratories (West Grove, PA, USA).

4.2.2. HSP90 enzymatic assay

HSP90 α Assay Kits (BPS Bioscience, San Diego, CA, USA) were utilized following the instruction manual, master mixture. HSP90 assay buffer, DTT, BSA, H₂O and FITC-labeled geldanamycin were added to all 96 wells. Then test compounds were added to each well designated "Test Inhibitor". To the "Blank", "Enzyme Positive Control" and "Enzyme Negative Control" wells, the same solution with inhibitor was added. HSP90 assay buffer was added to the "Enzyme Negative Control" and "Blank" wells. Finally, the recombinant protein HSP90 α was incubated in every well designated "Enzyme Positive Control" and "Test Inhibitor" for 2–3 h at rt with slow shaking. The fluorescent polarization of the samples was determined by microtiter-plate reader, which can detect excitation ranging from 475 to 495 nm and emission ranging from 518 to 538 nm.

4.2.3. SRB (sulforhodamine B) assay

Cells were seeded in 96-well plates and cultured overnight followed by the exposure to gradient concentrations of different compounds for 48 h. Briefly, cells in Tz group were fixed *in situ* with 10% trichloroacetic acid (TCA) to represent a measurement of the cell population at the time of drug addition (T₀). After an additional 48 h incubation with or without compounds in medium with 5% FBS, the assay was terminated by 10% TCA in CTL and treatment groups. SRB dye purchased from Sigma (St. Louis, MO, USA) at 0.4% (w/v) in 1% acetic acid was added to stain the cells. Unbound dye was removed by 1% acetic acid and the plates were air dried. Bound dye was subsequently solubilized with 10 mM trizma base, and the absorbance was read at a wavelength of 515 nm.

4.2.4. Colony formation assay

Cells were plated in 96 well plates (5×10^3 /well) and exposed to DMSO or compound **12c** at indicated concentrations for 24 h. The drugs were then washed away and cells were trypsinized and seeded in 24-well plates for continuing growth for 10 days. The colonies were fixed and stained with crystal violet (0.5% in 70% EtOH) and the experiments were repeated at least twice.

4.2.5. FACScan Flow Cytometric analysis

Cells were seeded in 6-well plates (2.5×10^5 /well) and treated with DMSO or **12c** at various concentrations for indicated times. Cells were washed with phosphate-buffered saline, fixed in ice-cold 70% EtOH at -20 °C overnight, and stained with propidium iodide (80 µg/ml) containing Triton X-100 (0.1%, v/v) and RNase A (100 µg/ml) in phosphate-buffered saline (PBS). DNA content was analyzed with the FACScan and CellQuest software (Becton Dickinson, Mountain View, CA, USA).

4.2.6. Immunoblotting and lentivirus expression system

Cells were seeded in dishes and allowed to attach overnight. The cells were treated with drugs at indicated concentrations for indicated times. After the indicated exposure time, cells were lysed and the immunoblotting was performed as previously described [24].

4.2.7. Statistics and data analysis

Each experiment was performed at least three times, and representative data are shown. Data in bar graphs are given as the means \pm S.D. Means were checked for statistical difference using the *t*-test and *P*-values less than 0.05 were considered significant (**P* < 0.05, ***P* < 0.01, ****P* < 0.001).

4.3. Molecular docking study

The crystal structure of HSP90 (PDB ID: 5GGZ) was obtained from Protein Data Bank [25]. Protein preparation included removal of water molecules from the crystal structure. Docking was performed using the Knime [26] software and the FlexX node (https:// www.biosolveit.de/knime/). The co-crystal structure was used to define the binding site. The 3D coordinates of docked compounds were generated using the "Generate 3D Coordinate" mode. Finally, all scoring parameters were used with the default settings.

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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Appendix A. Supplementary data

Supplementary data to this article can be found online at https://doi.org/10.1016/j.ejmech.2020.112540.

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