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Letter

Novel 2-Carbonylbenzo[b]thiophene 1,1-Dioxide Derivatives as Potent Inhibitors of STAT3 Signaling Pathway

Peng Ji,[†] Xin Xu,[‡] Shuhua Ma,[§] Junchao Fan,[†] Qiang Zhou,[†] Xinliang Mao,^{*,‡} and Chunhua Qiao^{*,†,‡}

[†]Department of Medicinal Chemistry, College of Pharmaceutical Science, Soochow University, 199 Ren Ai Road, Suzhou 215123, P. R. China

[‡]Jiangsu Key Laboratory of Translational Research and Therapy for Neuro-psycho-diseases, Department of Pharmacology, College of Pharmaceutical Sciences, and Jiangsu Key Laboratory of Preventive and Translational Medicine for Geriatric Diseases, Soochow University, Suzhou 215123, P. R. China

[§]Department of Chemistry, Jess and Mildred Fisher College of Science and Mathematics, Towson University, 8000 York Road, Towson, Maryland 21252, United States

(5) Supporting Information



ABSTRACT: Signal transducer and activator of transcription 3 (STAT3) is considered to be an attractive therapeutic target for cancer therapy. In this study, a series of 2-carbonylbenzo[b]thiophene 1,1-dioxide derivatives (**CBT**) were designed to inhibit the STAT3 SH2 domain phosphorylation site Try 705. We demonstrated that incorporation of basic flexible groups through amide bond linkage to benzo[b]thiophene 1,1-dioxide (**BTP**) achieved compounds with higher antiproliferative potency than **BTP** itself. The most potent compound **60**, as indicated from luciferase reporter gene assay, inhibited the STAT3 pathway by decreasing the phosphorylation level of STAT3 Tyr705, while the phosphorylation level of other upstream tyrosine kinases in this pathway was not significantly inhibited. Compound **60** was also shown to trigger ROS generation and accumulation, thus consequently attributed partially to the observed cell apoptosis. This study provided important structural information for the development of inhibitors targeting the STAT3 pathway.

KEYWORDS: STAT3, inhibitor, 2-carbonylbenzo[b]thiophene 1,1-dioxide derivatives, antiproliferative activity, apoptosis, reactive oxygen species

C ignal transducer and activator of transcription 3 (STAT3), \bigcirc one of seven different STAT isoforms (STAT1-6), regulates many critical functions in human normal and malignant tissues, such as cell proliferation, differentiation, apoptosis, immune response, and angiogenesis.¹⁻⁵ Upon binding to membrane-bound receptors, the STATs are brought into close proximity for the phosphorylation of the critical tyrosine residue (Tyr 705), the phosphorylated STAT3 undergoes dimerization through a reciprocal pTyr-SH2 domain interaction to form transcriptionally active STAT:STAT dimer, which translocates into the nucleus and binds to specific DNAresponse elements to promote STAT3 target gene transcription.⁶ Aberrant activation of STAT3 is reported in many human solid and hematological tumors, including ovarian, breast, colon, pancreatic, lung, brain, renal, and prostate cancers.⁷⁻⁹ On account of its crucial role in malignant transformation and tumorigenesis, STAT3 has emerged as a promising tumor therapeutic target.^{10–15}

The first reported small molecule STAT3 inhibitor is **STA21** (Figure 1), which is discovered to inhibit STAT3-dependent luciferase activity in breast and ovarian cancer cell lines.¹⁶ Derived from **STA21**, compound **LLL12** (Figure 1) also shows inhibitory effect on STAT3 phosphorylation, STAT3-DNA binding, and STAT3-dependent transcriptional activity.^{17–19} In addition, the substituted 1,4-naphthoquinone-based compound **Ly5** and **PD15** have been revealed to inhibit STAT3 pTyr705 phosphorylation.²⁰ Another structurally different skeleton like **Stattic** (Figure 1) is discovered as a STAT3 SH2 domain inhibitor, which selectively suppresses STAT3 protein activation, dimerization, and nuclear translocation in numerous cancer cell lines.²¹ Structurally similar to **Stattic**, compound **HJC0146** was also discovered to be a STAT3 signaling

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Figure 1. Representative STAT3 small molecule inhibitors and the design of compounds.

pathway include: **Carbazole**,²³ **Niclosamide**,^{24,25} **S3I-201**,^{26,27} **NSC-743380**,^{28,29} and so on. Most of the reported STAT3 inhibitors are demonstrated to target the pTyr-SH2 domain site Tyr705. Despite these reported efforts, none of these STAT3 inhibitors have been developed into a clinical trial, most possibly due to the lack of ideal physicochemical properties of the studied molecules.¹¹

To develop selective and potent STAT3 inhibitors, the benzo [b] thiophene 1,1-dioxide, as in **Stattic**, was employed as a leading scaffold. Previous studies have shown that this compound binds to the cleft of the STAT3 SH2 domain (PDB code 1BG1),²¹ which primarily consists of three sits: the polar and basic pTyr705 site, the adjacent hydrophobic side pocket, and the Leu706 site.^{30,31} By molecular docking analysis (Figure S1), it was envisioned that attachment of aromatic/ aliphatic moiety at 2-position through different chemical tethers (linker), such as amide, carbonyl, and ester (Figure 1), would enforce the BTP-protein interaction and consequently improve the compound biological potency and binding selectivity. For these reasons, a series of 2-carbonylbenzo [b] thiophene 1.1dioxide derivatives (CBT) were designed and synthesized, and their antiproliferative activity as well as the inhibitory effect toward the STAT3 pathway were studied. In addition, the biological effect of Stattic was shown to be associated with the formation of reactive oxygen species (ROS).^{32,33} Therefore, the CBT were evaluated on their ability to induce ROS generation. Collectively, this study led to the identification of CBT-based potent inhibitors targeting the STAT3 pathway.

Treatment of the substituted 2-fluorobenzaldehyde (1a-1d) with methyl ethyl 2-mercaptoacetate and potassium carbonate in *N*,*N*-dimethylformamide(DMF) at 60 °C resulted in the formation of methyl benzo[*b*]thiophene-2-carboxylate (2a-2d), and those were conveniently hydrolyzed to provide the corresponding acids 3a-3d (Scheme 1). Using 1-ethyl-3-(3dimethyllaminopropyl)carbodiiehydrochlide (EDC) as coupling reagent, the acids (3a-3d) or benzoic acid) were reacted with either aromatic amines or aliphatic amines to afford intermediates benzo[*b*]thiophene-2-carboxanilide (4a-4k, 4t, 4u). Compound *p*-tolyl 4-bromobenzo[*b*]thiophene-2-carboxylate (4l) was prepared by coupling of 4-bromobenzo[*b*]thiophene-2-carboxylic acid (3a) with *p*-cresol in the presence of DCC and DMAP. Intermediates 4m-4s and 4v were prepared from 3b and 3d with the corresponding secondary



1a, 2a, 3a, R_1 =Br, R_2 =H, R_3 =H; **1b, 2b, 3b**, R_1 =H, R_2 =Br, R_3 =H **1c, 2c, 3c**, R_1 =H, R_2 =H, R_3 =Br; **1d, 2d, 3d,** R_1 =H, R_2 =NO₂, R_3 =H 4a - 4s. 4v 6u 4a. 5a 5 4t, 4u 4t, 5a, R₄=phenyl 4u, 5v, R₄=piperidin-1-yl R₃=H, R₄=phenylamino **4I, 6I,** R₁=Br, R₂=H, R₃=H, R₄=*p*-tolyloxy 6a, R₁=Br, R₂=H **4b, 6b,** R₁=Br, R₂=H, R₃=H, R₄=*p*-tolylamino **4c, 6c,** R₁=Br, R₂=H, R₃=H, R₄=(4-chlorophenyl)amino **4d, 6d,** R₁=Br, R₂=H, R₃=H, R₄=(3-chlorophenyl)amino n, R₁=H, R₂=Br, R₃=H, R₄=diethylamino **4n, 6n,** R_1 =H, R_2 =Br, R_3 =H, R_4 =pyrrolidin-1-yl **4o, 6o,** R_1 =H, R_2 =Br, R_3 =H, R_4 =piperidin-1-yl 40, 60, R,=Er, R_2=H, R_3=H, R_4=(3-chlorophenyl); 46, 66, R,=H, R_2=BR, R_3=H, R_4=phenylamino 47, 67, R_1=H, R_2=H, R_3=Br, R_2=phenylamino 49, 69, R_1=H, R_2=Br, R_3=H, R_4=phenylamino 41, 61, R_1=H, R_2=Br, R_3=H, R_4=tort-butylamino 41, 61, R_1=H, R_2=Br, R_3=H, R_4=tort-butylamino 44, 64, R_1=H, R_2=Br, R_3=H, R_4=cyclohexylamino 44, 64, R_1=H, R_2=Br, R_3=H, R_4=cyclohexylamino 4p, 6p, R₁=H, R₂=Br, R₃=H, R₄=2-methylpip eridin-1-v 4a. 6a. R₁=H.R =H,R₄=3-methylpiperidin-1-y $\begin{array}{l} \mathbf{y}_{1}, \mathbf{y}_{1}, \mathbf{y}_{1}, \mathbf{y}_{1}, \mathbf{y}_{2}, \mathbf{y}_{3}, \mathbf{z}_{1}, \mathbf{x}_{4} = 3, \text{methylpiperidin-1-yl} \\ \mathbf{4r}, \mathbf{6r}, \mathbf{7}, \mathbf{4r}, \mathbf{R}, \mathbf{R}, \mathbf{R}, \mathbf{R}, \mathbf{2r}, \mathbf{R}, \mathbf{3} = \mathbf{R}, \mathbf{R}, \mathbf{3r} = \mathbf{R}, \mathbf{0} \\ \mathbf{4r}, \mathbf{6r}, \mathbf{R}, \mathbf{1}, \mathbf{R}, \mathbf{R}, \mathbf{2r}, \mathbf{R}, \mathbf{3r} = \mathbf{R}, \mathbf{R}, \mathbf{1}, \mathbf{R}, \mathbf{1}, \mathbf{R}, \mathbf{2r} = \mathbf{0} \\ \mathbf{4r}, \mathbf{6r}, \mathbf{R}, \mathbf{1}, \mathbf{R}, \mathbf{R}, \mathbf{2r} = \mathbf{0} \\ \mathbf{4u}, \mathbf{6u}, \mathbf{R}, \mathbf{1}, \mathbf{R}, \mathbf{R}, \mathbf{2r} = \mathbf{0} \\ \mathbf{4u}, \mathbf{6u}, \mathbf{R}, \mathbf{1}, \mathbf{R}, \mathbf{R}, \mathbf{2r} = \mathbf{0} \\ \mathbf{4r}, \mathbf{6r}, \mathbf{R}, \mathbf{1}, \mathbf{R}, \mathbf{R}, \mathbf{2r} = \mathbf{0} \\ \mathbf{4r}, \mathbf{R}, \mathbf{1}, \mathbf{R}, \mathbf{$

"Reagents and conditions: (a) SHCH₂COOCH₃, K₂CO₃, DMF, 60 °C; (b) KOH, MeOH, 60 °C; (c) (i) for (4a-4k, 4t-4u) RNH₂, EDC, pyridine, RT; (ii) for (4l) *p*-cresol, DCC, DMAP, DCM, RT; (iii) for (4m-4s, 4v) secondary amines, HOBt, EDC, TEA, DCM, RT, overnight; (d) Pd/C, H₂, MeOH; (e) *m*-CPBA, DCM, 45 °C.

amines using EDC and 1-hydroxybenzotriazole (HOBt) as coupling reagents. Treatment of 4g and 4v with Pd/C gave compounds 5a and 5v, which were coupled with benzoic acid to give 4t and 4u. Finally, all benzo[b]thiophenes were oxidized to 2-carbonylbenzo[b]thiophene 1,1-dioxides using *m*-chloroperbenzoic acid (*m*-CPBA) as the oxidant.

We first evaluated the antiproliferative activity of these compounds against selected cancer cell lines, which are reported with overexpression of STAT3 protein: breast cancer cell lines include MCF-7 (ER-positive), MDA-MB-231 (ERnegative and triple-negative), and MDA-MB-435S (ER-negative and triple-negative), and nonbreast cancer cell lines include the prostate cancer cell line DU145, the pancreatic cancer cell line PANC-1, and the nonsmall cell lung cancer cell line A549. The established MTT assay was used to determine the compound IC₅₀ values with Stattic as positive control. As shown in Table 1, except 6l, all prepared compounds exhibited significant antiproliferative activity against most tested cancer cells, with $IC_{50}s$ in single digit micromole range. For compound **61**, the *p*cresol has an ester linkage with 4-bromobenzo[b]thiophene-2carboxylic acid and showed the weakest inhibitory activity. All other compounds with the amide linkage ensured their good antiproliferative activity. Apparently, a variety of aliphatic amines at the position of R_4 were well tolerated (6i, 6j, and 6m–6s). These compounds exhibited higher activity against all selected cancer cells. This result was consistent with our molecular docking analysis. By comparison, aromatic amines at R_4 resulted in derivatives with relatively low potency (6a-6g and 6t). In addition, large size substituents at R_2 or R_4 exerted unfavorable influence on the potency of compounds (6g, 6k, 6t, and **6u**).

To demonstrate whether or not these compounds would show the expected selectivity between normal cell vs cancer cell, the IC_{50} s of compounds **6a**, **6j**, and **6o** against normal human liver cell HL-7702 and liver cancer cell line SMMC-

Table 1. Effect of Newly Synthesized Analogues on the Proliferation of Human Cancer Cell Lines

		$\mathrm{IC}_{50}^{b}(\mu\mathrm{M}) \pm \mathrm{SEM}$					
		breast cancer cells			nonbreast cancer cells		
compd	cLogP ^a	MDA-MB-231	MDA-MB-435S	MCF-7	DU145	PANC-1	A549
6a	3.01	1.74 ± 0.53	1.39 ± 0.19	1.73 ± 0.15	2.64 ± 0.15	6.67 ± 0.27	4.09 ± 0.32
6b	3.19	2.46 ± 0.95	n.d. ^{<i>c</i>}	n.d.	3.03 ± 0.05	n.d.	n.d.
6c	3.83	3.78 ± 0.89	n.d.	n.d.	6.34 ± 0.21	n.d.	n.d.
6d	3.85	2.21 ± 0.77	n.d.	n.d.	3.39 ± 0.60	n.d.	n.d.
6e	3.03	1.72 ± 0.39	n.d.	n.d.	2.77 ± 0.30	n.d.	n.d.
6f	3.00	1.84 ± 0.47	n.d.	n.d.	2.66 ± 1.08	n.d.	n.d.
6g	1.90	3.04 ± 0.60	n.d.	n.d.	5.19 ± 1.12	n.d.	n.d.
6h	2.49	2.31 ± 0.07	n.d.	n.d.	3.86 ± 0.68	n.d.	n.d.
6i	2.49	1.34 ± 0.36	0.82 ± 0.08	1.27 ± 0.16	1.35 ± 0.01	3.34 ± 0.15	2.85 ± 0.71
6j	2.28	1.09 ± 0.21	0.93 ± 0.04	1.24 ± 0.10	2.17 ± 0.23	5.05 ± 1.68	2.97 ± 0.26
6k	3.11	5.73 ± 0.68	n.d.	n.d.	5.73 ± 0.50	n.d.	n.d.
61	3.69	>15	n.d.	n.d.	>15	n.d.	n.d.
6m	2.45	1.67 ± 0.03	1.54 ± 0.19	0.89 ± 0.11	1.46 ± 0.09	1.94 ± 0.08	1.58 ± 0.07
6n	1.71	1.27 ± 0.41	0.94 ± 0.12	1.38 ± 0.29	1.54 ± 0.14	3.47 ± 0.24	3.47 ± 1.25
60	2.05	0.70 ± 0.34	0.79 ± 0.09	0.91 ± 0.07	1.03 ± 0.29	2.81 ± 0.46	1.70 ± 0.29
6р	2.48	0.95 ± 0.21	0.98 ± 0.11	0.99 ± 0.05	0.98 ± 0.11	2.14 ± 0.16	1.3 ± 0.08
6q	2.67	1.07 ± 0.42	1.37 ± 0.17	1.26 ± 0.16	1.41 ± 0.26	2.98 ± 0.15	1.94 ± 0.16
6r	1.44	1.04 ± 0.49	0.80 ± 0.14	1.05 ± 0.16	1.91 ± 0.26	3.95 ± 1.34	2.31 ± 0.45
6s	2.85	0.88 ± 0.12	0.95 ± 0.04	1.13 ± 0.09	1.26 ± 0.14	4.38 ± 0.78	2.08 ± 0.05
6t	3.30	3.21 ± 0.50	n.d.	n.d.	6.21 ± 0.79	n.d.	n.d.
6u	2.64	1.68 ± 0.90	n.d.	n.d.	2.07 ± 0.42	n.d.	n.d.
Stattic	0.90	1.56 ± 0.56	1.87 ± 0.11	2.16 ± 0.29	2.50 ± 0.31	2.90 ± 0.56	2.50 ± 0.37

^{*a*} cLogP: http://146.107.217.178/lab/alogps/start.html. ^{*b*} The inhibitory effects of individual compounds on the proliferation of cancer cell lines were determined by the MTT assay. The data are the mean \pm SD from at least three independent experiments. ^{*c*}n.d.: not determined (for compounds with IC₅₀ values higher than **Stattic** against MDA-MB-321 and DU145 cells, their antiproliferative activity was not determined in other cancer cells).

7721 were determined. As presented in Figure 2A, their IC₅₀ values were 2.4–6.0 μ M against liver cancer cell line SMMC-7721. By comparison, the IC₅₀s against normal human liver cell line HL-7702 were 15–25 μ M, suggesting that these compounds exhibited high toxicity and good selectivity for cancer cells.

Based on the IC₅₀ values in Table 1, compound **60** was selected to investigate the mechanism of inhibition by using STAT3-dependent dual luciferase reporter assay in HeLa cells, which expresses a high level of constitutive STAT3. As shown in Figure 2B, **60** inhibited STAT3 activity in a concentration-dependent manner. Compound **60** at 2 μ M decreased 14% of the STAT3-induced luciferase activity; when the concentration of **60** was doubled to 4 μ M, the luciferase activity was decreased by 48%, which was higher than the positive STAT3 inhibitor **Stattic**. At the same concentration, **Stattic** was decreased by 36%. These results indicated that **60** displayed a remarkable inhibitory effect on STAT3-induced luciferase activity in HeLa cells.

The effect of **60** on the intracellular STAT3 activation was further investigated using Western blot analysis. The MDA-MB-231 breast cancer cells were treated with **60** for 12 h, the total cell lysates were prepared and the Y705 phosphorylated STAT3 proteins were detected using specific antibody. As shown in Figure 2C–E, phosphorylation of Tyr705 was inhibited by **60** in a dose-dependent manner, while the total level of STAT3 protein was not affected at these concentrations. Moreover, **60** displayed higher inhibitory effect on STAT3 phosphorylation than **Stattic** in all the tested cells (Figure 2C–E). By contrast, **60** did not show marked inhibition on the phosphorylation of other kinases including Src, Erk1/2, and mTOR (Figure 3A), which are all involved in



Figure 2. (A) Compounds **6a**, **6j**, and **6o** showed lower toxicity against normal human liver cells HL-7702 than liver cancer cells. (B) Compound **6o** inhibited the STAT3-mediated luciferase activity in transfected HeLa cells. (A,B) Data were expressed as mean \pm SD from experiments performed in triplicate. ((*) p < 0.05, (**) p < 0.01, (***) p < 0.001 compared with vehicle-treated control). (C–E) Compound **6o** affected the STAT3-Y705 phosphorylation in MDA-MB-231, MDA-MB-435S, and MCF-7 cells. (F) **6o** inhibited IL-6-induced STAT3 phosphorylation in the MDA-MB-231 cell.

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Figure 3. (A) Compound **60** had no obvious effect on upstream tyrosine kinases. (B) Compound **60** increased the cleaved PARP levels in MDA-MB-231 in a dose-dependent manner. (C) Compound **60** increased the cleaved PARP levels in MDA-MB-231 in a time-dependent manner. (D) Compound **60** up-regulated the cleavage of PARP in three breast cancer cells. (E) Apoptotic ratio of different concentrations of **60** in MDA-MB-231 and MCF-7 cells. The values are presented as means \pm SD from at least three independent experiments ((*) p < 0.005, (**) p < 0.01, (***) p < 0.001 compared with vehicle-treated control).

the STAT3 signaling pathway. These results thus indicated that **60** could effectively inhibit the STAT3 phosphorylation in a manner probably independent of these associated kinases (Figure S2).

To further evaluate the inhibition of **60** on STAT3 activity, MDA-MB-231 and MCF-7 cells were starved overnight, followed by treatment with **60** for 2 h before IL-6 stimulation. As shown in Figure 2F, IL-6 markedly increased the phosphorylation level of STAT3, which was decreased by **60** in a concentration-dependent manner. Compound **60** at 4 μ M almost completely abolished STAT3 phosphorylation induced by IL-6. These results revealed that **60** effectively inhibited IL-6-induced STAT3 phosphorylation (Figure S3).

To investigate whether **60** could induce apoptosis, the poly(ADP-ribose) polymerase (PARP) was used as a characteristic indication to demonstrate the early apoptotic event in the intact cells. After treatment of MDA-231 cells with **60**, cleavage of PARP protein was determined by Western blot analysis. Notably, **60** elevated the cleaved PARP levels in a time- and concentration-dependent manner (Figure 3B,C). In addition, cleavage of PARP was also observed in both MDA-MB-435S and MCF-7 cells with **60** treatment (Figure 3D).

The percentages of apoptotic cells were further determined by flow cytometry. As shown in Figure 3E, the percentage of the apoptotic cells increased in a concentration-dependent manner after treatment with compound **60**. In the MDA-MB-231 cells, these values were 10.5%, 32.7%, and 63.1% (early and late apoptosis) at the indicated concentration, while only 4.8% was observed in the control cells. Similarly, the induced apoptosis rates in MCF-7 at 0, 2, 4, and 6 μ M were 2.6%, 13.9%, 45.0%, and 75.8%, respectively. These results suggested that compound **60** could efficiently induce cancer cell apoptosis, which partially explained the antiproliferative effect of **60** on human cancer cells (Figure S4).

It was reported that benzo[b]thiophenesulphonamide 1,1dioxide derivatives could induce reactive oxygen species (ROS)-mediated apoptotic process in tumor cells.³² To investigate whether or not our newly synthesized derivatives could also induce ROS production, MDA-MB-231 and MCF-7 cells were pretreated with **60**, and then cells were collected and stained by DCFH-DA and subjected to flow cytometry analysis. As shown in Figure 4A, **60** triggered a dose-dependent ROS



Figure 4. (A) Measurement of ROS production in MDA-MB-231 and MCF-7 cells by using a ROS-sensitive fluorometric probe. (B) Cleaved-PARP was determined by Western blot analysis ((*) p < 0.05, (**) p < 0.01, (***) p < 0.001 compared with vehicle-treated control).

generation in both MDA-MB-231 and MCF-7 cancer cells, which would partially contribute to the observed PARP protein cleavage and cell apoptosis.

Next, Western blot analysis was conducted to detect whether the **60**-induced cleavage of PARP could be abrogated by addition of a ROS scavenger. The MDA-MB-231 cancer cells were pretreated with an antioxidant, *N*-acetylcysteine (NAC). Then, compound **60** was added, and PARP was collected and determined. As depicted by Figure 4B, the antioxidant NAC could significantly decrease **60**-induced cleavage of PARP. Besides, **60** could decrease the intracellular GSH content in the MDA-MB-231, which confirmed **60** induced apoptosis partially by interacting with GSH (Figure S5). These results verified our hypothesis that compound **60** could induce the generation and accumulation of ROS, which was possibly involved in the cell apoptosis by interacting with GSH.

In summary, a series of 2-carbonylbenzo[b]thiophene 1,1dioxide derivatives (**CBT**) were developed to inhibit the phosphorylation of the STAT3. Through incorporation of basic flexible groups to the benzo[b]thiophene 1,1-dioxide (**BTP**) core structure, higher potency compounds were discovered. We confirmed that the compounds inhibited the phosphorylation of STAT3 and cell proliferation and induced the apoptotic process of cancer cells with constitutively active STAT3. The present study provided more structural reference for the development of a useful tool in elucidating the inhibition of STAT signaling pathway. Further *in vivo* antitumor activity studies on these compounds are undergoing and will be reported in due course.

ASSOCIATED CONTENT

Supporting Information

The Supporting Information is available free of charge on the ACS Publications website at DOI: 10.1021/acsmedchem-lett.5b00228.

Supplemental data, spectroscopic data with methods of synthesis of all compounds, and methods of biological evaluation (PDF)

AUTHOR INFORMATION

Corresponding Authors

*E-mail: xinliangmao@suda.edu.cn.

*E-mail: qiaochunhua@suda.edu.cn.

Notes

The authors declare no competing financial interest.

ABBREVIATIONS

DCM, dichloromethane; DCC, dicyclohexylcarbodiimide; DMAP, 4-dimethylaminopyridine; TEA, triethylamine; RT, room temperature; SD, standard deviation; MTT, 3-(4,5dimethyl-2-thiazolyl)-2,5-diphenyl-2-*H*-tetrazolium bromide; DCFH-DA, 2',7'-dichlorofluorescin diacetate

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