



Recent updates on thienopyrimidine derivatives as anticancer agents

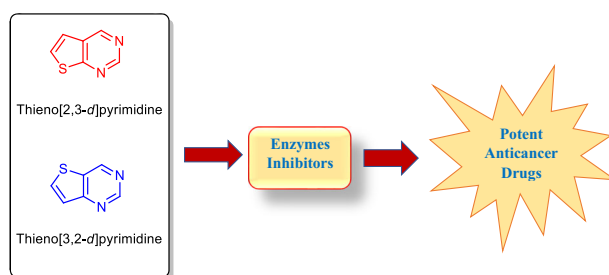
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Received: 4 November 2022 / Accepted: 13 February 2023 / Published online: 2 March 2023
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Abstract

Thienopyrimidine derivatives hold a unique place between fused pyrimidine compounds. They are important and widely represented in medicinal chemistry as they are structural analogs of purines. Thienopyrimidine derivatives have various biological activities. The current review discusses different synthetic methods for the preparation of heterocyclic thienopyrimidine derivatives. It also highlights the most recent research on the anticancer effects of thienopyrimidines through the inhibition of various enzymes and pathways, which was published within the last 9 years.

Graphical Abstract



Keywords Thienopyrimidine · Synthesis · Anticancer agents · Enzyme inhibition

Introduction

Cancer is the most significant health problem with increasing frequency and mortality rates globally. Due to the large number of cancer patients, effective treatment approaches and prompt diagnosis are imperative [1]. Cell proliferation and differentiation of cancer cells are originated from the action of some enzymes including protein kinases (PKs) [2]. As a result of mutations in PKs, oncogenesis can occur, and these mutations are critical to the progression of cancer [3]. As a result, the use

of PK inhibitors has become increasingly important in the last two decades since PKs are one of the pathways that can be inhibited in cancer treatment to solve a variety of cellular communication problems [4]. In clinical oncology, PKs are frequently used as molecular therapeutic targets because they play key roles in several signal transduction pathways, which can lead to metastasis and drug resistance. [5]. Developing kinase inhibitors as anticancer medicines continue to be a crucial research priority to improve tumor selectivity, efficiency, and safety of anticancer medicines. Furthermore, there are other targets that can be inhibited to give effective anticancer drugs such as topoisomerases [6, 7], tubulin polymerization [8], and histone deacetylase (HDAC) [9]. Thienopyrimidine scaffold is one of the most frequently used chemical scaffolds in drug development. The structural and isoelectronic characteristics of thienopyrimidine-containing compounds are similar to those of purine and they have

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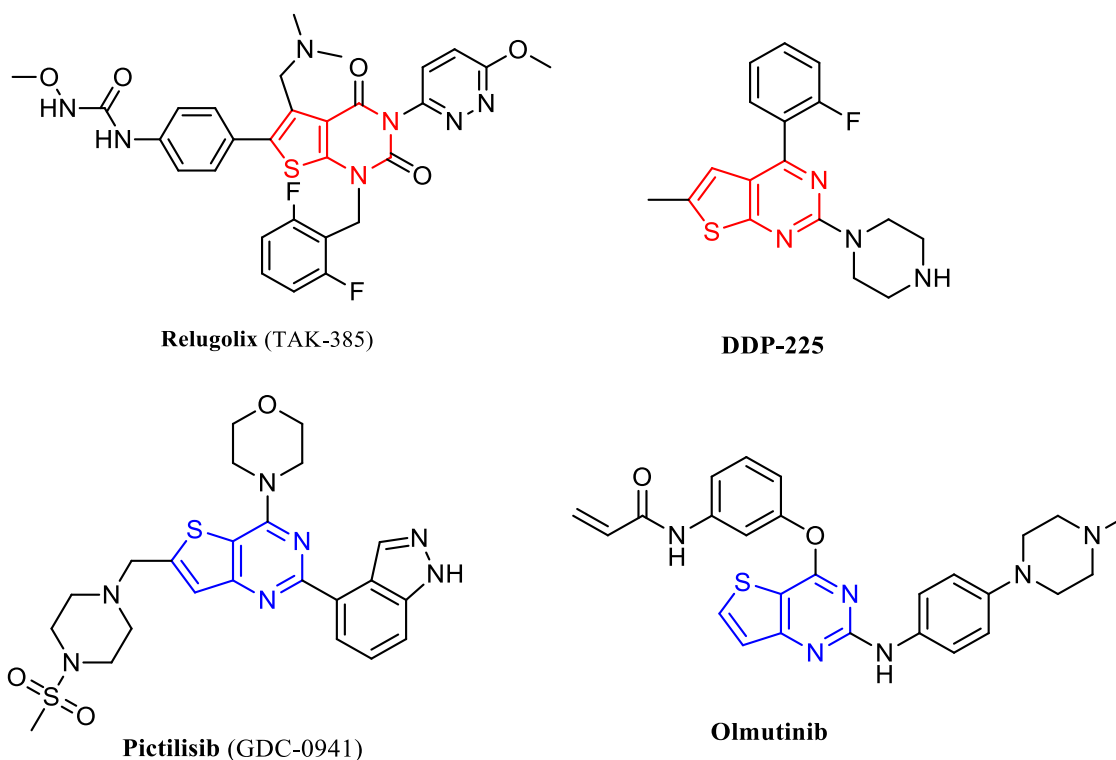


Fig. 1 Structures of some thienopyrimidine-containing drugs

become an attractive structural feature in the production of pharmaceutical drugs [10, 11]. Thienopyrimidines have been demonstrated to have significant and various pharmacological properties, such as antibacterial [12–14], antiviral [15, 16], anti-inflammatory [17, 18], antiprotozoal [19], and anticancer activities [20–23]. Figure 1 represents some thienopyrimidine-containing drugs with varying profiles of biological activity. Relugolix (TAK-385), is a thienopyrimidine derivative that has completed phase III clinical trials and is being studied for its capacity to treat endometriosis and prostate carcinoma by acting as a gonadotropin-releasing hormone receptor (GnRHR) antagonist [24, 25]. DDP-225 is another thienopyrimidine drug that entered phase II clinical trials and was designed to cure irritable bowel syndrome (IBS) and gastrointestinal tract (GIT) diseases by acting as a serotonin receptor (5-HT₃) antagonist and noradrenaline reuptake inhibitor [26, 27]. Moreover, pictilisib (GDC-0941) is a thieno[3,2-d]pyrimidine derivative which inhibits phosphatidylinositol 3-kinase (PI3K) and is in clinical trials and was clinically investigated for the treatment of advanced solid tumors [28]. In addition, olmutinib is a marketed drug that inhibits epidermal growth factor receptor (EGFR) and is used to treat NSC lung cancer [29, 30].

Synthetic strategy

In the literature, several synthetic pathways are reported that involve the construction of either the pyrimidine ring or the

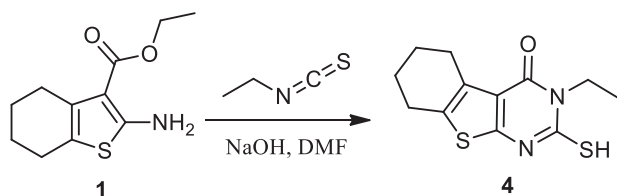
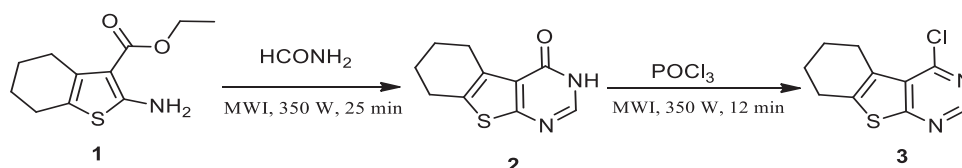
thiophene ring to obtain the polysubstituted thienopyrimidines. Thienopyrimidines have been prepared using 2-amino-3-substituted thiophene derivatives as important starting compounds. Gewald's procedure was used in the traditional synthesis of these derivatives [31–33]. The synthesis of thieno[2,3-*d*]pyrimidine scaffolds has been described using a variety of methods [34–37]. Our goal is to report the synthesis of thienopyrimidine derivatives using two main strategies. Either pyrimidine ring closure in aminothiophene derivatives or thiophene ring closure in pyrimidine derivatives.

Starting from thiophene derivatives

Starting from 2-aminothiophene-3-carboxylate derivatives

Phoujdar et al. reported a microwave-based synthesis of novel thienopyrimidine derivatives that were designed as gefitinib bioisosteres in high yield [38]. The synthesis of compound **2** via the reaction of 2-aminoester derivative **1** with formamide using microwave irradiation (MWI) took 25 minutes, whereas the previous traditional method took from 8–10 hours [39] (Scheme 1). Moreover, chloro derivative **3** was produced when compound **2** reacted with phosphorus oxychloride under MWI for 12 minutes. In addition to improved yield and purity, the reaction time for this method was reduced from 14 h to 12 min.

Moreover, Mavrova et al. synthesized thiosemicarbazide and 1,3,4-thiadiazole thieno[2,3-*d*]pyrimidine derivatives

Scheme 1 Synthesis of thieno[2,3-*d*]pyrimidines **2** and **3****Scheme 2** Synthesis of thieno[2,3-*d*]pyrimidine **4**

[40] (Scheme 2). The formation of the pyrimidine ring of compound **4** was achieved by cyclocondensation of the 2-aminoester derivative **1** with ethyl isothiocyanate in presence of NaOH.

Starting from amino cyanothiophene derivatives

In 2014, Kerru et al. synthesized derivatives of thienopyrimidines that contain 1,2,4-triazoles and 1,3,4-oxadiazoles [41] (Scheme 3). Compound **6** was obtained by refluxing 5-amino-4-cyanothiophene derivatives **5** and triethyl orthoformate. Moreover, triazolo derivatives **7** were produced from the reaction of compound **6** with substituted aryl hydrazides in toluene under reflux.

On the other hand, Gao et al. demonstrated that thienopyrimidine derivative **9** was produced by the reaction of substituted 2-aminothiophene-3-carbonitrile **8** with trifluoroacetic acid (TFA) in presence of toluene and phosphorus oxychloride [42] (Scheme 4).

Starting from 2-aminothiophene-3-carboxamide derivatives

Kassab et al. synthesized a series of hexahydrocyclooctathieno[2,3-*d*]pyrimidines [43] (Scheme 5). Cyclocondensation of 2-aminothiophene-3-carboxamide derivative **10** with aromatic aldehydes in dry dimethylformamide provided thienopyrimidine derivatives **11**. Refluxing derivatives **11** and phosphorus pentasulfide in presence of xylene, produced 4-thioxo derivatives **12**.

Meanwhile, Rashad et al. synthesized thienopyrimidine derivatives [44] (Scheme 6). Compound **13** produced the equivalent 2-thioxo derivative **14** when heated with carbon disulfide under reflux.

Starting from pyrimidine derivatives

Thienopyrimidines could be synthesized from pyrimidine derivatives. Brough et al. reported the production of

compound **16** by the reaction of compound **15** with ethyl-2-mercaptoacetate in presence of potassium carbonate as a base. Upon reaction of the ester **16** with aqueous ammonia and microwave irradiation at 130 °C, compound **17** was produced [45] (Scheme 7).

Saddik et al. produced thieno[2,3-*d*]pyrimidine derivatives in 2018 [46] (Scheme 8). Compound **18** reacted with thiourea in ethanol, followed by treatment with sodium hydroxide solution and acidification with diluted HCl, to yield 4-mercapto-2-morpholino-6-phenylpyrimidine-5-carbonitrile **19**. Moreover, compound **19** undergoes cyclization to give derivatives **20** through alkylation with chloroacetonitrile, chloroacetamide, and ethyl chloroacetate in ethanol and in the presence of potassium carbonate.

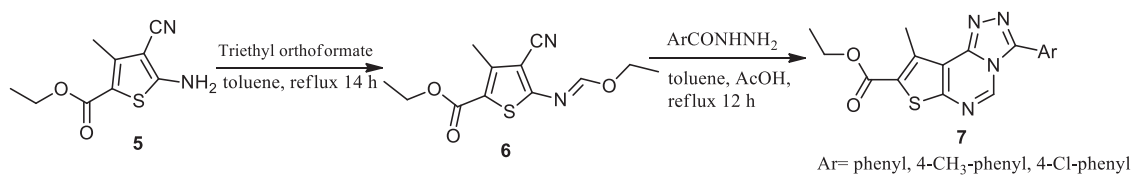
Anticancer activity

Nowadays cancer is the most dangerous life-threatening disease in our life. It is suggested to be the first reason for mortality in the future. The number of cancer patients globally is supposed to increase during the next years [47]. Biological studies of thieno[2,3-*d*]pyrimidines have demonstrated that the replacement of different groups on this important core confers antineoplastic activity via inhibition of various kinases [48–50].

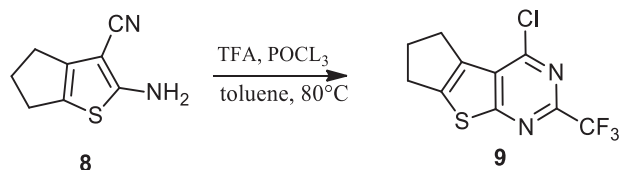
Thienopyrimidine derivatives as protein kinase inhibitors

Thienopyrimidine derivatives as epidermal growth factor receptor (EGFR) inhibitors

In small molecular cancer therapy, epidermal growth factor receptor tyrosine kinase (EGFR TK) is an important target [51, 52]. It is a cell-surface tyrosine kinase receptor that is stimulated by the alpha transforming growth factor (TGF α), extracellular protein ligands, and members of the epidermal growth factor (EGF) family [53]. EGFR overexpression has been linked to uncontrolled cell division in a variety of cancers, including multiform glioblastoma, lung and anal carcinoma [54]. In 2014, Yang et al. produced a series of thienopyrimidine derivatives with α,β -unsaturated amide side chains at position 6 (Fig. 2) [55]. Compound **21** was of great interest because it was found to be better than the marketed drug lapatinib as an EGFR inhibitor. Moreover, it showed



Scheme 3 Synthesis of thieno[2,3-*d*]pyrimidines **7**



Scheme 4 Synthesis of thieno[2,3-*d*]pyrimidine **9**

better activity than lapatinib against breast carcinoma (SK-BR-3) cell line with $IC_{50} = 0.13 \mu\text{M}$. It displayed irreversible inhibition of the EGFR enzyme due to the existence of an amide side chain that creates a covalent bond with Cys773 placed in the ATP pocket of the EGFR enzyme [55].

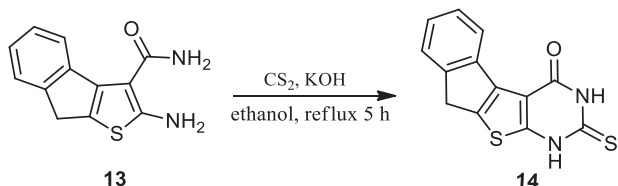
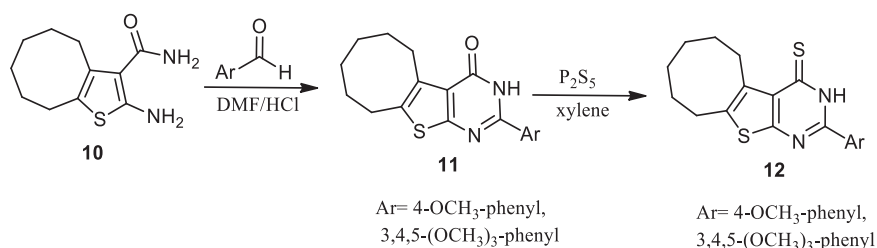
A novel series of 6-cinnamoyl-4-arylaminothiopyrimidines was synthesized in 2020. They were evaluated as anticancer agents and displayed highly potent cytotoxic activity in comparison to erlotinib (Fig. 2) [56]. Thieno[2,3-*d*]pyrimidine derivatives **22a-h** and thieno[3,2-*d*]pyrimidine derivatives **23a-f** were created. The anti-neoplastic activity of these compounds against prostate cancer (PC3) showed that all of the compounds demonstrated excellent activity with IC_{50} values in the sub-micromolar values from 0.1 to $0.79 \mu\text{M}$. All of the derivatives **23a-f** demonstrated a significant effect on prostate cancer PC3; breast cancer MDA-MB-231 and hepatocellular cancer cell line (HepG2) with IC_{50} from 0.10 to $15.90 \mu\text{M}$, and moderate activity on lung cancer cell line (A549) with IC_{50} from 6.67 to $26.24 \mu\text{M}$.

In the two series **22a-h** and **23a-f**, the presence of an ethynyl group connected to the aryl amine group at C-3 resulted in good cytotoxic effects on PC3 and breast cancer (MDA-MB-231) cell lines when compared to other cell lines. 3,5-Dichloro substituted derivative **22e** was the most potent derivative against all of the examined cell lines, followed by derivative **22g** with 3,4-dichloro substitution. The most powerful compounds **22e** and **22g** were evaluated on colorectal cancer (HCT-116 and SW480); breast cancer (SKBR3); ovarian cancer (SKOV3) and glioblastoma cell (U87) cell lines. They demonstrated IC_{50} ranging from 3.83 to $11.94 \mu\text{M}$, compared to erlotinib which exhibited lower effectiveness against these cell lines with IC_{50} from 22.99 to $61.78 \mu\text{M}$. In addition, western blot analysis revealed that compound **22e** inhibited the phosphorylation of EGFR and downstream molecule ERK1/2. Furthermore, the effects of compounds **22e** and **22g** on cell cycle distribution and

apoptosis were investigated, and the results revealed that most cells remained in the G0 phase and that cell growth was arrested. Moreover, it was revealed that for **22e** and **22g**, the percentages of early and late apoptosis were 11%, 15.5%, and 42.8%, 15.7%, correspondingly so they stimulated cell death. Molecular docking study showed that compound **22e** formed hydrogen binding interactions with Asp 831, Met 769, and Lys 721. In addition to π - π and π -alkyl interactions with Phe 699 and Val 702, respectively in the hydrophobic pocket. Moreover, in the solvent region, the cinnamamide part created Van der Waals interactions with Gly 772 and Pro 770 (Fig. 3). From this study we can summarize the presence of dichloro substitution on aniline ring may enhance the anticancer activity and the cinnamamide moiety is favorable to interact with the active site of EGFR enzyme.

Thienopyrimidine derivatives as vascular endothelial growth factor receptor 2 (VEGFR-2) inhibitors

The VEGF receptors are a kind of receptor tyrosine kinase (RTK) that is important for vascular development and hematopoiesis [57]. The three VEGFR members are VEGFR-1, VEGFR-2, and VEGFR-3 [58]. VEGFR performs a crucial starring role in the proliferation, migration, and angiogenesis of vascular endothelial cells when it is activated by VEGF [59, 60]. The significance of VEGFR-2 in tumor angiogenesis has prompted attention to the progress of VEGFR-2 inhibitors [61–63]. Sorafenib [64], sunitinib [65, 66], lenvatinib [67], and linifanib [68] are examples of VEGFR-2 inhibitors. In 2015, Abdel Aziz et al. synthesized tricyclic pyrido[3',2':4,5]thieno[3,2-*d*]pyrimidin-4-amine derivatives as VEGFR-2 inhibitors (Fig. 4) [69]. It was found that compound **24a** with thienopyridine ring demonstrated the strongest inhibition against VEGFR-2 by 67% with an IC_{50} value of $2.6 \mu\text{M}$, whereas compound **24b** with pyrazolopyridine and **24c** with isoxazolepyridine inhibited VEGFR-2 by 12% and 18%, respectively. Additionally, the molecular docking of **24a** revealed that Cys917 in the adenine region of the ATP binding site generated the crucial hydrogen bonding connection with its core structure and hydrophobic interactions between the 4,6-dimethylthieno[2,3-*b*]pyridine group and Val897, Cys1043, and Leu1033 at the other end of the ATP binding site. Moreover, 7,9-dimethyl substituted

Scheme 5 Synthesis of thieno[2,3-*d*]pyrimidines **11** and **12****Scheme 6** Synthesis of thieno[2,3-*d*]pyrimidine **14**

pyrido[3',2':4,5]thieno[3,2-*d*]pyrimidine was participated in interactions with Ala864 and Val914 (Fig. 5). From the previous findings, we conclude that the existence of thieno[3,2-*d*]pyrimidine as a core structure is important to interact with the ATP binding site of VEGFR-2.

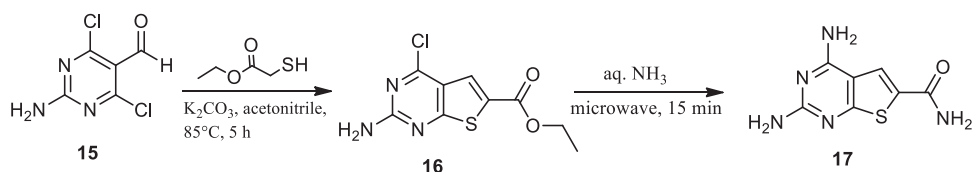
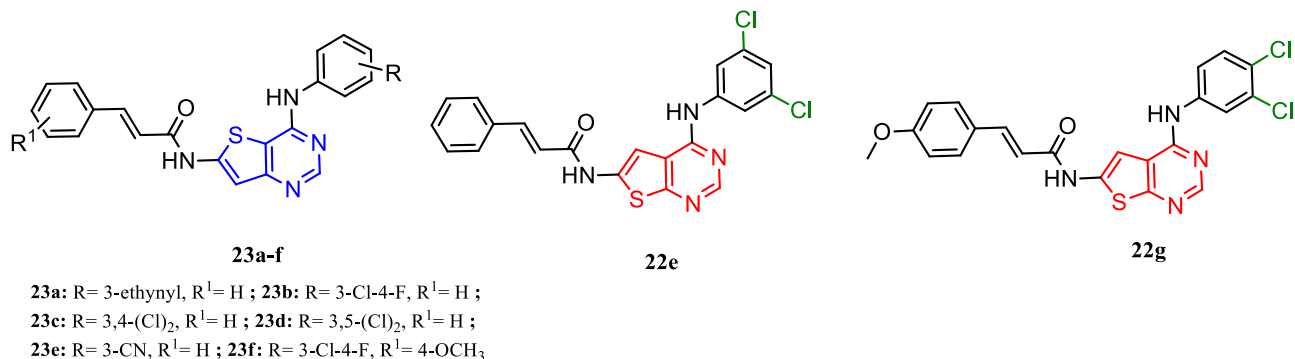
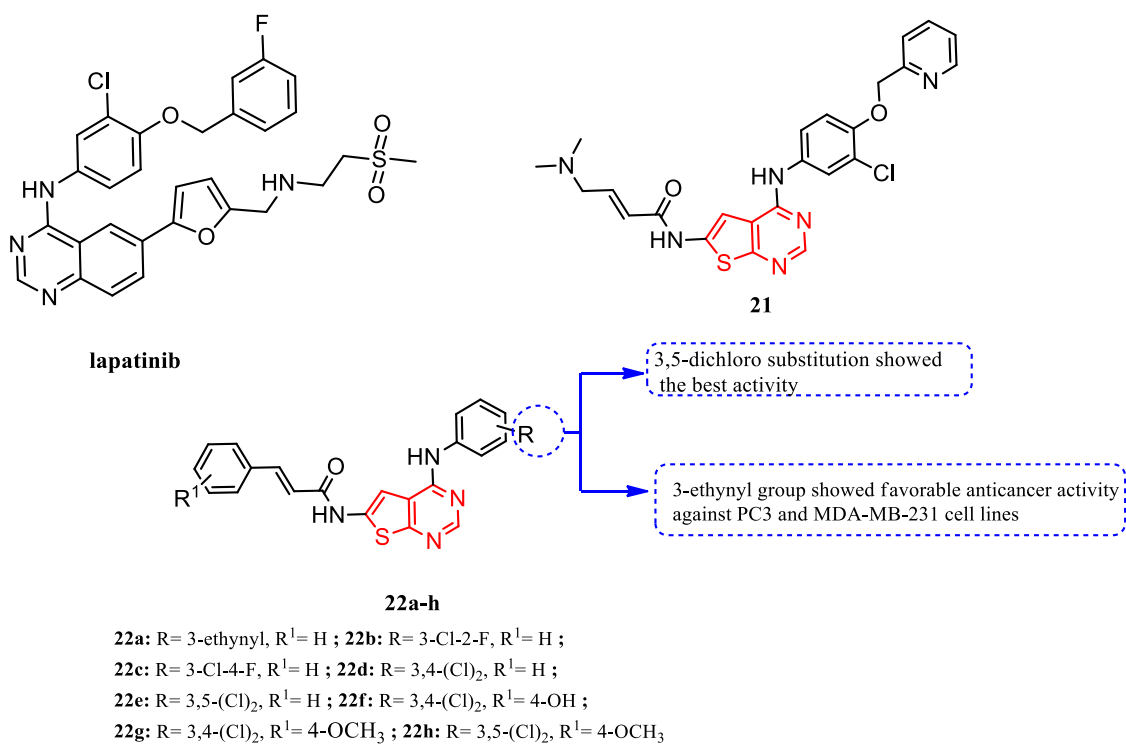
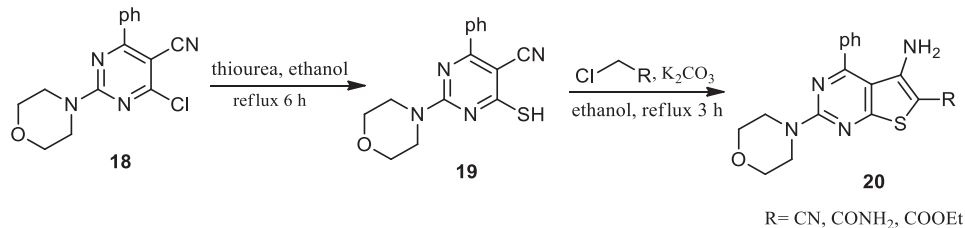
In 2021, new hybrid compounds of thieno[2,3-*d*]pyrimidine with aryl aminothiazole were designed and evaluated as VEGFR-2 kinase inhibitors (Fig. 4) [70]. In relation to the synthesized compounds, it was discovered that the addition of a weak electron-withdrawing halogen atom, such as 4-chloro, 3-bromo, or 4-bromo, results in more effective carcinogenic agents. However, the compound with unsubstituted phenyl **25a** and the compound substituted with a strong electron withdrawing as the 3-nitro group **25b** displayed little cytotoxic activity against all investigated cell lines. Moreover, it was found that the replacement of 4-chloro atom in compound **25c** with a 4-bromo atom in compound **25e**, decreased the anticancer activity with mean inhibition of 18.30% and 4.35%, correspondingly. Additionally, the position of the bromo group in **25e** was changed from para to meta in **25d**, which resulted in an increase in the mean inhibition from 4.35% (**25e**) to 16.83% (**25d**). Therefore, the 4-chloro derivative **25c** showed highly powerful VEGFR-2 kinase inhibition with an IC₅₀ value of 62.48 nM in comparison to sorafenib. Additionally, compound **25c** was the most effective derivative versus CNS cancer (SNB-75 and SF-295), and renal cancer (CAKI-1) cell lines with IC₅₀ values of 7.12 ± 0.33, 7.36 ± 0.39, and 4.84 ± 0.22 μM, respectively. Furthermore, the results of the flow cytometric study demonstrated that **25c** exhibited cytotoxic activity by inhibiting cellular growth and causing cell cycle arrest at the G2/M phase. Moreover, molecular modeling showed the ability of compound **25c** to form interactions with essential amino acids in the VEGFR-2 binding site as demonstrated in Fig. 5.

From the previous results, we can assume the substitution of terminal phenyl ring with halogen is more beneficial in anticancer activity than substitution with a strong electron-withdrawing group.

Moreover, in 2021, new derivatives of thieno[2,3-*d*]pyrimidines were synthesized as VEGFR-2 kinase inhibitors [71]. Among the synthesized compounds, **26a-d** and **27** showed the most potent inhibition against VEGFR-2 kinase with IC₅₀ values ranging from 0.23 ± 0.03 to 0.37 ± 0.04 μM. Additionally, compound **26b** with 4-chlorophenyl showed the most powerful VEGFR-2 kinase inhibition with an IC₅₀ value of 0.23 ± 0.03 μM. For compound **26b**, it was found to have greater anticancer activity than sorafenib against colorectal carcinoma HCT-116 and hepatocellular carcinoma HepG2 cell lines with IC₅₀ of 2.80 ± 0.16 and 4.10 ± 0.45 μM, respectively. It was found that the replacement of an electron-withdrawing group (4-Cl) in **26b** with an electron donating group (4-OCH₃) in **26f**, resulted in a loss of activity. However, compound **26e** with a 2-methoxyphenyl group showed enhanced biological activity with VEGFR-2 inhibition IC₅₀ of 0.69 ± 0.06 μM. Molecular docking of compound **26b** revealed that hydrogen bonds were created between the hydrazide group and Glu883 and Asp1044. In addition, Cys1043 and Val897 formed two hydrophobic interactions with the phenyl ring (spacer). Moreover, 5,6,7,8-tetrahydrobenzo[4,5]thieno[2,3-*d*]pyrimidine moiety created hydrophobic interactions with Leu1033, Leu838, Ala864, Cys917 and Val846 and a hydrogen bond with Cys917. Finally, the terminal phenyl ring interacted with Ile886 through a hydrophobic bond (Fig. 5). From the previous study, we can adopt the presence of thieno[2,3-*d*]pyrimidine as a core structure is important to interact with the ATP binding site of VEGFR-2 and the substitution of terminal phenyl ring with electron-withdrawing group is more useful in anticancer activity than substitution with electron donating group.

Thienopyrimidines derivatives as PI3K/AKT/mTOR pathway inhibitors

The phosphatidylinositol 3-kinase (PI3K)/ protein kinase B (AKT)/ mammalian target of rapamycin (mTOR) (PI3K/AKT/mTOR) signaling system stands as a major mechanism that controls cell existence, proliferation, glucose

Scheme 7 Synthesis of thieno[2,3-*d*]pyrimidine **17****Scheme 8** Synthesis of thieno[2,3-*d*]pyrimidines **20****Fig. 2** SAR of thienopyrimidines as EGFR inhibitors

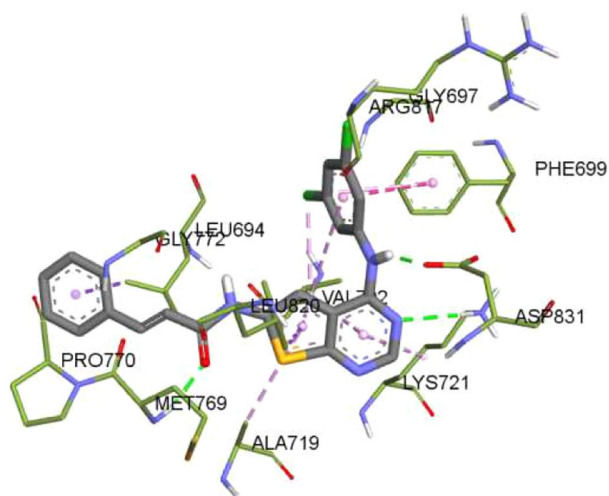


Fig. 3 Interactions of compound **22e** inside EGFR binding site (PDB: 1M17)

metabolism, migration, and death [72, 73]. Throughout the last few decades, it has been widely explored to develop new cancer therapies. There are now several medications that target this route in clinical studies, the most predominant of which is class **I** PI3K inhibitors [74, 75]. In 2021, Sun et al. synthesized and evaluated a novel series of thieno[2,3-*d*]pyrimidine derivatives as PI3K inhibitors [76]. Compound **28** (Fig. 6) with unsubstituted morpholine moiety showed extraordinary PI3K inhibition with an IC_{50} value equal to 7.2 nM. In addition to good pharmacokinetics properties, and significant anticancer activity against gastric cancer cell line HGC-27 with IC_{50} of 0.39 μ M. Compound **28** formed a hydrogen bond with Val851 in the hinge region of the PI3K active site. Additionally, compound **28** created a hydrogen bond network with Tyr836, Asp801, and the conserved water molecule in the affinity pocket. Also, Lys802 and deprotonated sulfonamide exhibited a charged interaction (Fig. 7).

Recently in 2022, Elmenier et al. synthesized and evaluated a series of 2-aryl-4-morpholinothieno[2,3-*d*]pyrimidine derivatives as PI3K inhibitors against various isomers PI3K α , β , and γ in addition to their anticancer activity versus NCI 60 cell lines (Fig. 6) [77]. The enzymatic activity of compounds **29** and **30** with a 3-hydroxyphenyl ring was good for PI3K β (62% and 72%) and PI3K γ (70% and 84%), correspondingly. Furthermore, derivatives containing tetramethylene substitution at positions 5 and 6 of the thienopyrimidine **30** and **32** mainly revealed improved activity contrasted to 5-methyl-6-carboxylate derivatives **29** and **31**. Moreover, compounds **31** and **32** also showed reduced inhibitory activity when the hydroxyl position was changed from 3 to 4 with PI3K β % inhibition (39% and 50%) and PI3K γ % inhibition (33% and 36%), respectively. Therefore, compound **30** was

demonstrated as the strongest inhibitor (72% and 84% on PI3K β and PI3K γ , respectively). Additionally, a molecular docking study of compound **30** with PI3K β showed that morpholine moiety formed a hydrogen bond with Val848 and the 3-hydroxyl group formed two hydrogen bond interactions with Lys799 and Asp931. Moreover, the thienopyrimidine ring formed hydrophobic interaction with Met773. On the other hand, docking of **30** with PI3K γ demonstrated the formation of three hydrogen bonds with Val882, Asp964, and Asp841 and hydrophobic interaction between the tetramethylene ring and Met953. Furthermore, the morpholine moiety created hydrophobic interaction with Ile881 (Fig. 7). Finally, according to this study, maintaining the morpholine part that binds to Val residue at the hinge region is one of the most important factors to take into account in the design of an efficient PI3K inhibitor.

In 2020, Han et al. synthesized a new series of thieno[3,2-*d*]pyrimidine derivatives that contain aryl hydrazide part which were evaluated as PI3K/mTOR dual inhibitors (Fig. 6) [78]. This study showed that the aryl hydrazide on C-6 was most beneficial since the hydrazide moiety is strongly desired by the solvent-exposed region of PI3K α and the presence of methoxy group on the terminal phenyl ring enhanced the activity. Moreover, the inhibition activity of compounds substituted with indazole **33b** or 2-aminopyrimidine **33c** groups on C-2 was greater than the inhibition activity of morpholino-substituted derivative **33a**. Therefore, compound **33c** with a 4-methoxybenzohydrazide group at C-6 and 2-aminopyrimidine group at C-2 of the thieno[3,2-*d*]pyrimidine backbone demonstrated the most effective PI3K α and mTOR inhibitory activities with IC_{50} values of 0.46 and 12 nM, respectively. Also, **33c** inhibits PI3K γ with an IC_{50} value of 13 nM. Moreover, the cell cycle study of **33c** revealed cell cycle arrest in the G1/S stages, which caused the HCT-116 cells to undergo apoptosis. The docking study of compound **33c** with PI3K γ displayed that the oxygen atom of morpholine moiety formed a hydrogen bond with Val882 and the NH of aryl hydrazide formed two H-bonds with Asp950. Moreover, Thr 887 created a hydrogen bond with the carbonyl group of the aryl hydrazide. Additionally, at the terminal phenyl ring, a 4-methoxy group formed a hydrogen bond with Lys 890, while the 2-aminopyrimidine fragment at the C-2 formed hydrogen bonds with Asp 841 and Asp 964. When compound **33c** was docked in the mTOR active site, comparable hydrogen bond interactions were detected whereas the morpholine moiety formed a hydrogen bond with Val882, the 2-aminopyrimidine fragment at the C-2 also formed hydrogen bonds with Asp 964 and Asp 841 (Fig. 7). From the previous three studies, we can assume the importance of the presence of unsubstituted morpholine moiety while designing effective PI3K inhibitors as it binds to Val residue at the hinge region of PI3K active site.

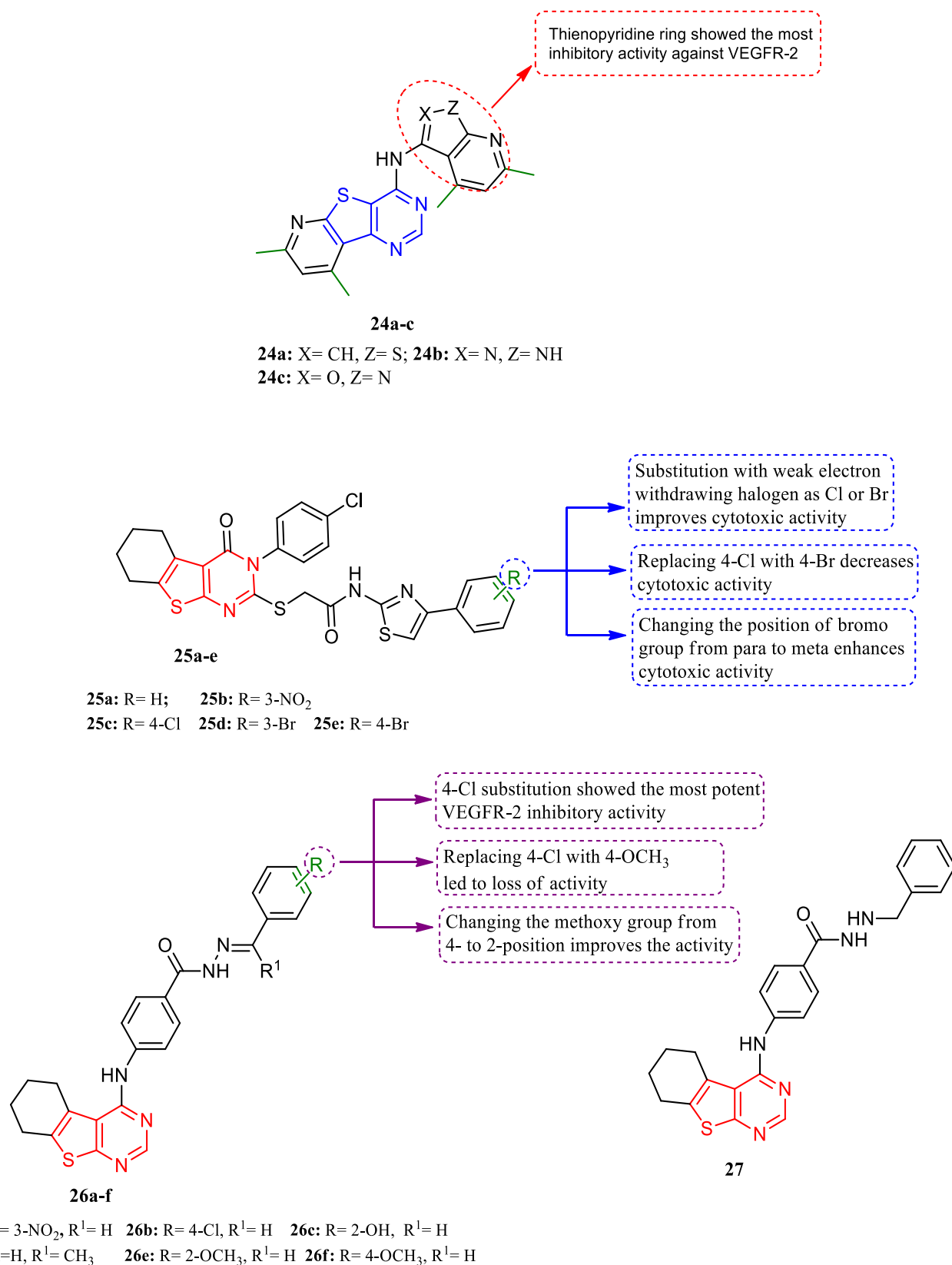


Fig. 4 SAR of thienopyrimidines as VEGFR-2 inhibitors

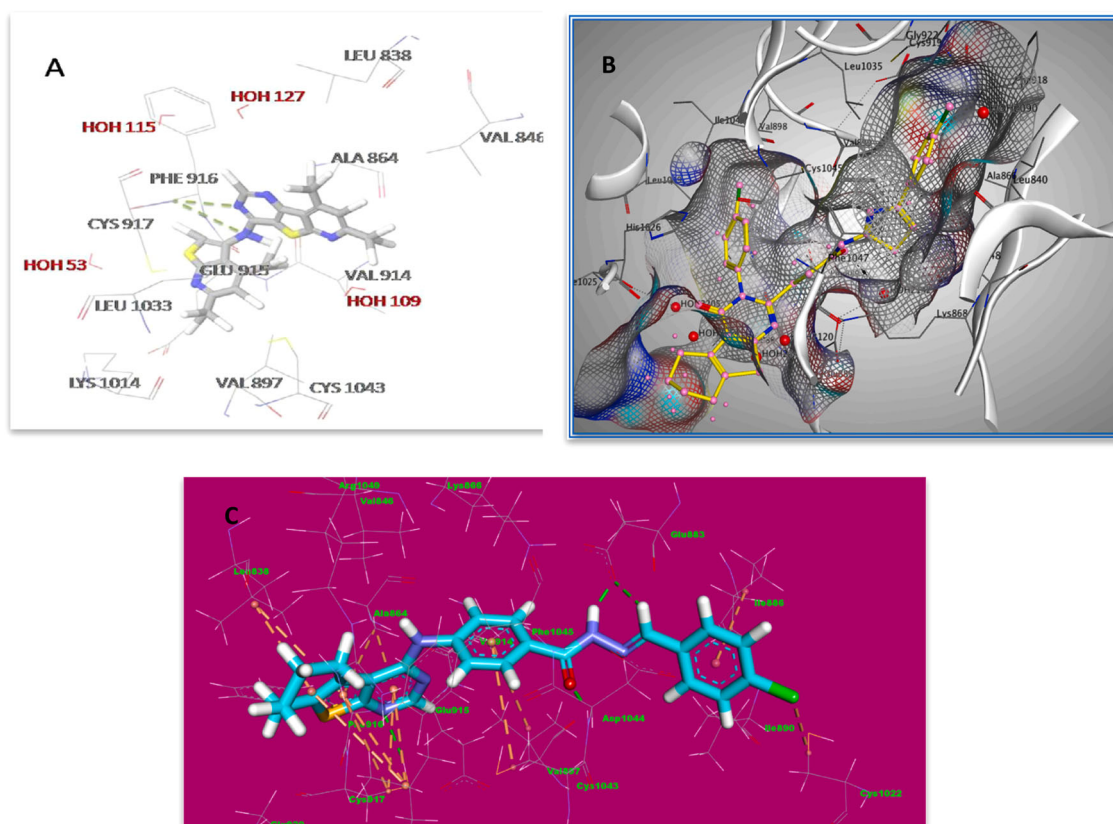


Fig. 5 **A** Interactions of compound **24a** inside VEGFR-2 binding site (PDB: 1YWN); **B** Interactions of compound **25c** inside VEGFR-2 binding site (PDB: 4ASD); **C** Interactions of compound **26b** inside VEGFR-2 binding site (PDB: 2OH4)

Thienopyrimidine derivatives as EGFR/ HER2 dual inhibitors

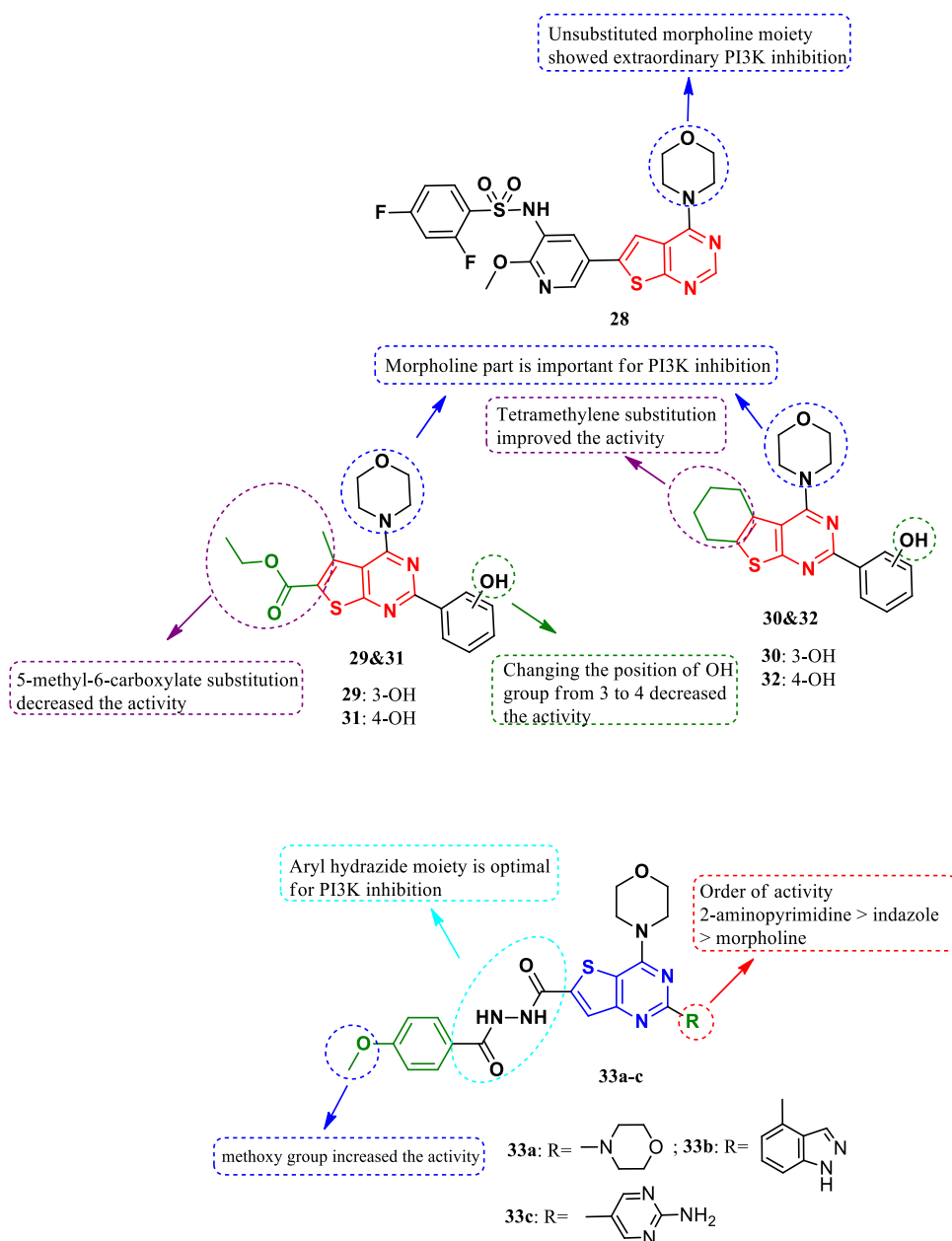
The ErbB family consists of four members: EGFR (ErbB-1), human epidermal growth factor receptors (HER2, ErbB-2), catalytically inactive ErbB-3, and ErbB-4 [79]. When a ligand interacts with the extracellular domain of either EGFR or HER2, kinase-active homodimers and heterodimers are formed [80, 81]. EGFR TK inhibitors have been successfully used as a treatment for NSC lung cancer patients [82]. The main disadvantage of administering EGFR TK inhibitors is the development of secondary or acquired resistance [83]. Because overactive EGFR and HER2 tyrosine kinases are essential hallmarks of various cancers, including colorectal, lung, pancreatic, head, and neck cancers, dual blocking of the EGFR and HER2 pathway is an excellent strategy for effective anticancer therapy [84–86].

In 2016, Abd El Hadi et al. reported the synthesis of two series of 4-anilinothieno[2,3-*d*] pyrimidines which were assessed as dual EGFR/HER2 kinase inhibitors (Fig. 8) [87]. Series **A** contains compounds that were less potent inhibitors for EGFR/HER2 kinase than Series **B**. Hence, the replacement of the 5-methyl-6-carboxylate groups in series **A** with the more hydrophobic 5,6-tetramethylene moiety in series **B** led to a significant increase in EGFR/HER2

inhibition. On the other hand, 3-chloroaniline-containing derivatives were more active than derivatives with *m*-unsubstituted aniline. Five derivatives of the tested compounds from series **B** (**34–38**) showed significant EGFR/HER2 inhibitory action, as measured by their IC_{50} values as demonstrated in Table 1. Therefore, compound **37** exhibited the highest inhibitory activity on both kinases. Regarding the molecular docking of compound **37** with EGFR active site, it showed hydrogen bond interaction between N1 of thienopyrimidine ring and Met793 and it formed a water-mediated hydrogen bond through N3 with Thr854. Additionally, **37** created π -cation and π -sigma interactions with Lys745 and Phe856, respectively. On the other hand, compound **37** demonstrated crucial interactions with HER2 binding site as N1 of thienopyrimidine created hydrogen bond interaction with Met801 as well as the formation of π -cation interaction with Lys753 (Fig. 9). From the mentioned work we conclude that, the presence of a more hydrophobic moiety attached to the thienopyrimidine ring is more effective towards EGFR/HER2 inhibition.

In 2018, Milik et al. prepared a series of thieno[2,3-*d*] pyrimidine as dual EGFR/HER2 kinase inhibitors built on the 6-phenylthieno[2,3-*d*]pyrimidine as a core scaffold (Fig. 8) [88]. Compounds **39a–c** provided potent dual

Fig. 6 SAR of thienopyrimidines as PI3K/AKT/mTOR inhibitors



EGFR/HER2 inhibitory activity with IC_{50} values of 21.4, 47.7, and 91.7 nM and 1.5, 0.879, and 1.2 μ M, respectively. This study showed that these compounds are equipped with bulky aniline head groups able to penetrate into the EGFR and HER2 back pockets. Moreover, compounds **39a-c** inhibited breast cancer SKBR3 cell line with IC_{50} equals 6.0, 4.7, and 4.83 μ M, respectively. Also, with IC_{50} of 4.2 μ M, **39c** greatly inhibited the growth of the non-small cell lung cancer NCI-H1975 cell line and enhanced the percentages of apoptotic and necrotic cells. Molecular docking and interactions of compound **39a** with key amino acids inside EGFR and HER2 active sites are presented in Fig. 9. We determine from the previous work the

significance of the presence of bulky aniline group at position 4 of the thienopyrimidine ring in designing EGFR and HER2 inhibitors.

Thienopyrimidine derivatives as VEGFR-2/ BRAF kinases dual inhibitors

Rapidly accelerated fibrosarcoma (RAF) kinases are serine/threonine protein kinases (PKs) that show a significant role in cell survival and proliferation. ARAF, BRAF, and CRAF are members of the RAF family whereas, in human malignancies, the BRAF valine 600 residue (V600E) mutation is the most common [89]. Melanomas, colorectal,

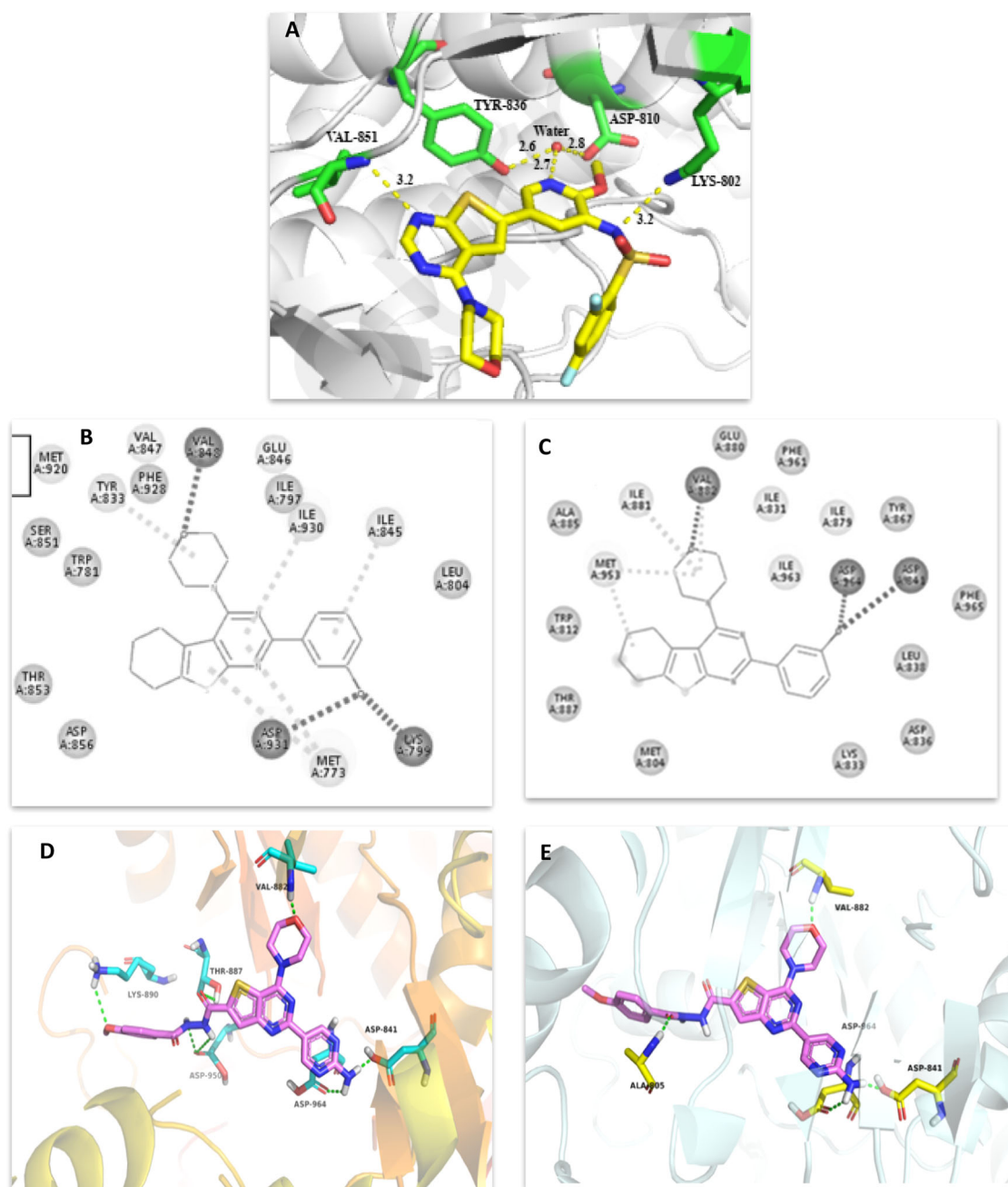


Fig. 7 **A** Interactions of compound **28** inside PI3K binding site (PDB: 4ZOP); **B** Interactions of compound **30** inside PI3K β binding site (PDB: 2Y3A); **C** Interactions of compound **30** inside PI3K γ binding site (PDB: 3DBS); **D** Interactions of compound **33c** inside PI3K γ binding site (PDB: 3DBS); **E** Interactions of compound **33c** inside mTOR binding site (PDB: 3L16)

thyroid cancer, and other human cancers can all be treated with BRAf inhibitors [90–92]. Consequently, dual inhibition of BRAF /VEGFR-2 is seen as a viable cancer therapeutic technique [93, 94]. Recently in 2022, Hassan et al. synthesized new hexahydrobenzo[4,5]thieno[2,3-*d*]pyrimidine derivatives as dual VEGFR-2/BRAF inhibitors (Fig. 10) [95]. Compounds **40** and **41** were elongated with the crucial urea insertion to have distal moieties, which increased antiproliferative action. Additionally, adding a *p*-chloro

group to the terminal phenyl ring of compound **41** slightly increased its antiproliferative activity as compared to derivative **40**. Therefore, compounds **40** and **41** demonstrated significant anticancer activity against most cancer cell lines. In addition, compounds **40** and **41** effectively inhibited VEGFR-2, BRAF^{V600E}, and BRAF^{WT} with IC₅₀ values of 0.111 ± 0.006 and 0.049 ± 0.003 μM, 0.089 ± 0.005 and 0.063 ± 0.003 μM, and 0.071 ± 0.004 and 0.05 ± 0.003 μM, respectively, compared to sorafenib. Furthermore,

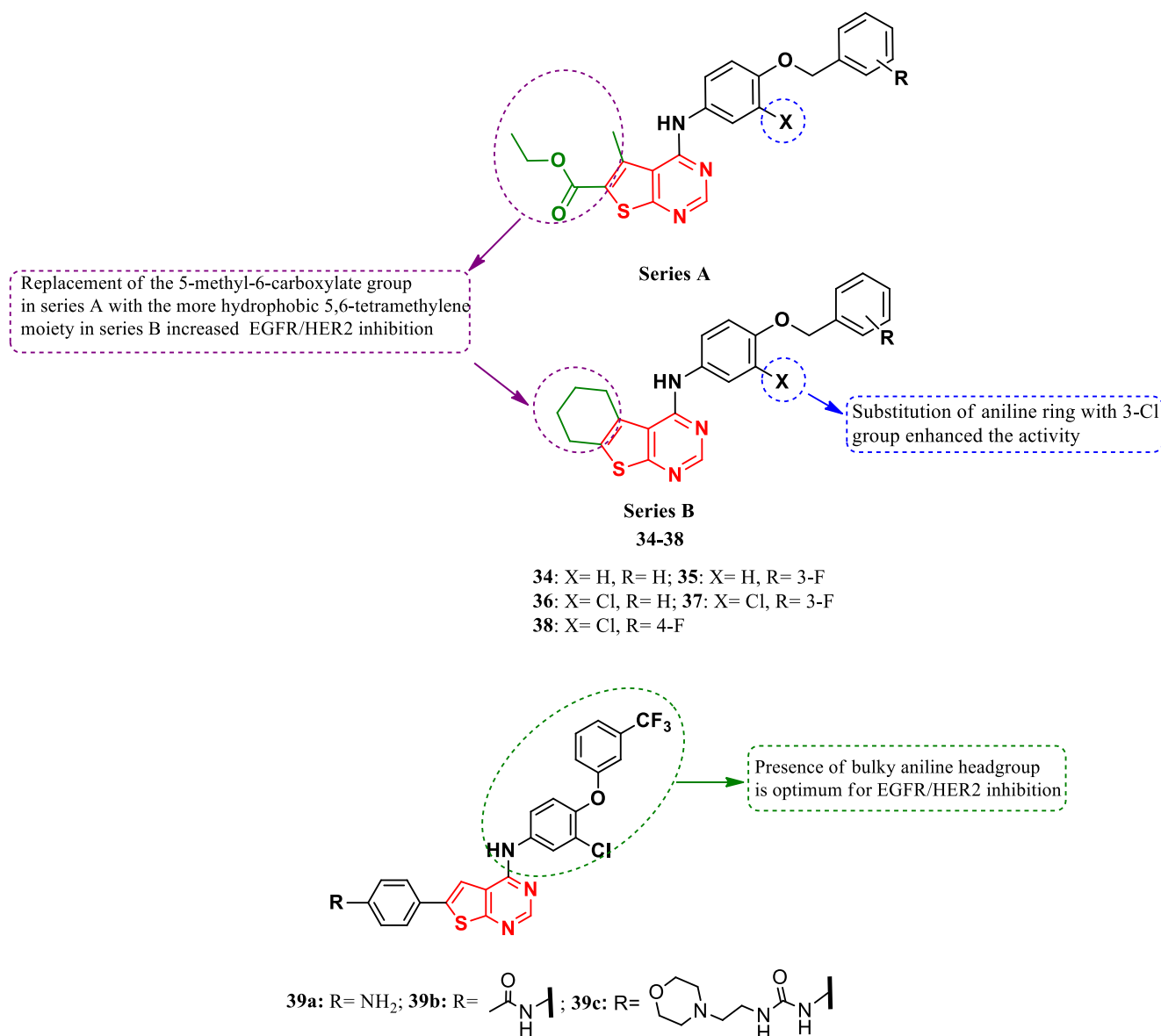


Fig. 8 SAR of thienopyrimidines as EGFR/HER2 dual inhibitors

Table 1 IC₅₀ (μM) of the most potent compounds of series B

Compound	X	R	EGFR/IC ₅₀ (μM)	HER2/IC ₅₀ (μM)
34	H	H	1.2	8.2
35	H	3-F	0.6	3.4
36	Cl	H	0.3	1.3
37	Cl	3-F	0.2	0.5
38	Cl	4-F	0.4	2.7

compounds **40** and **41** increased the overall apoptotic proportion in the breast cancer MCF7 cell line by 22.82 and 25.81 fold, correspondingly. Additionally, the examination of the cell cycle revealed that compounds **40** and **41** primarily arrested the cell cycle in the G1 and G1/S phases,

correspondingly. When compounds **40** and **41** docked inside VEGFR-2 and BRAF binding sites, they exhibited comparable interactions. Regarding VEGFR-2, 4-chlorophenyl cycloalkylthieno[2,3-*d*]pyrimidine formed a hydrogen bond with Cys919 through N1 as well as π - π and π -H interactions with Phe1047 and Leu840, respectively. Moreover, Glu885 and Asp1046 demonstrated hydrogen bond interactions with the urea group, and a sulfur-dipole interaction took place between the sulfur atom in 2-thioacetamide moiety and Glu917. Also, in **41** the chloro atom at the terminal phenyl group formed halogen bonding with Ile1025. On the other hand, docking of **40** and **41** inside BRAF showed hydrogen bond interactions between urea moiety with Glu500 and Asp593. Furthermore, the 2-thioacetamide spacer made a hydrogen bond with Thr528.

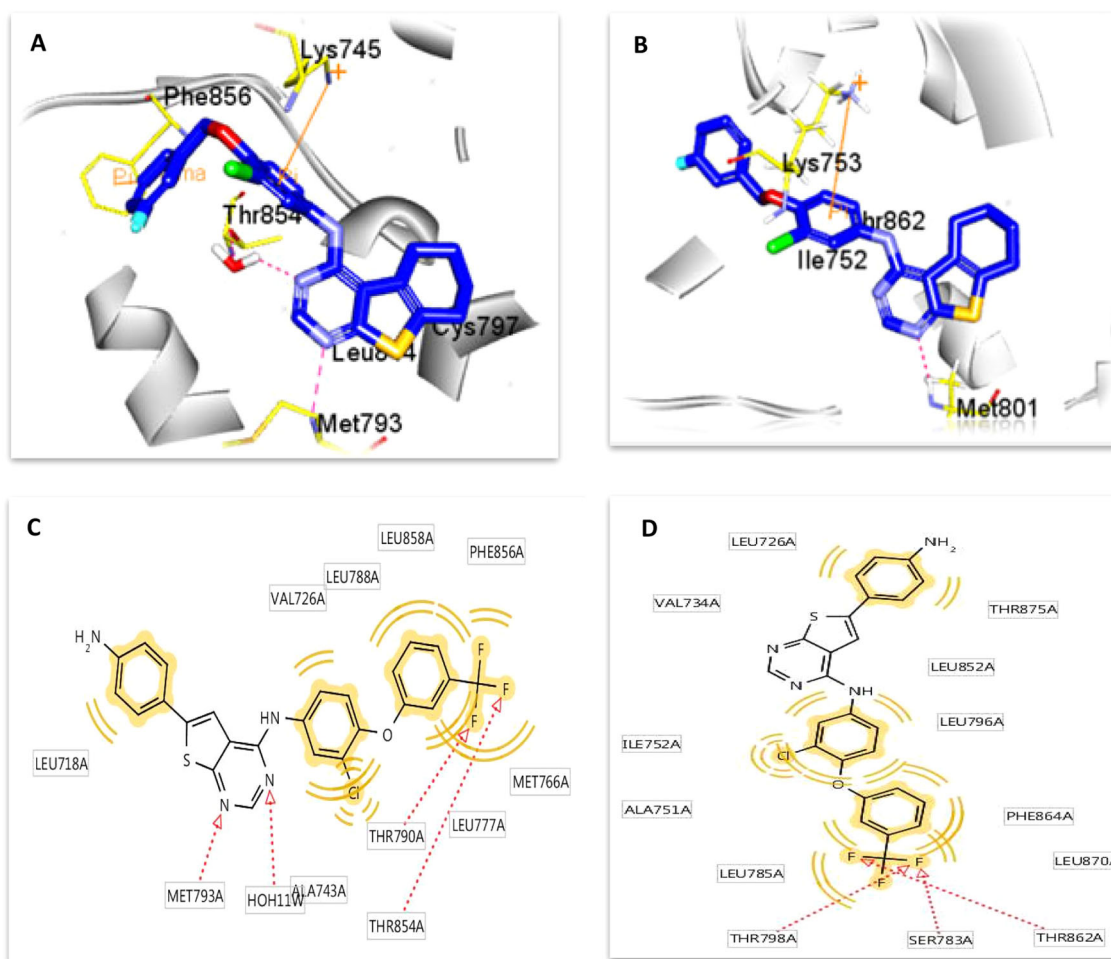


Fig. 9 **A** Interactions of compound **37** inside EGFR binding site (PDB: 1XKK); **B** Interactions of compound **37** inside HER2 binding site (PDB: 3RCD); **C** Interactions of compound **39a** inside EGFR binding

site (PDB: 3POZ); **D** Interactions of compound **39a** inside HER2 binding site (PDB: 3RCD)

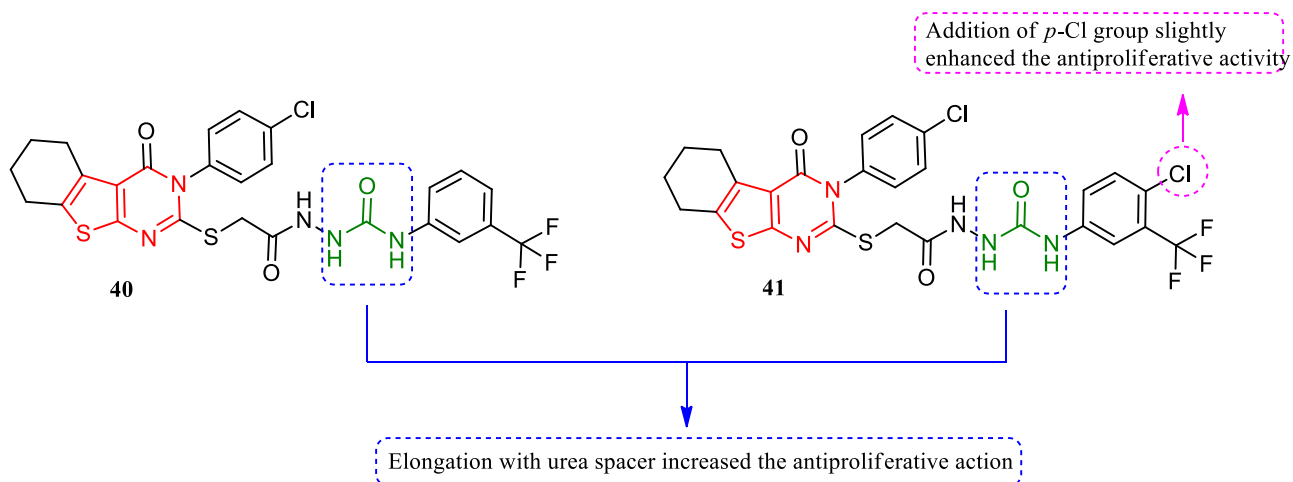
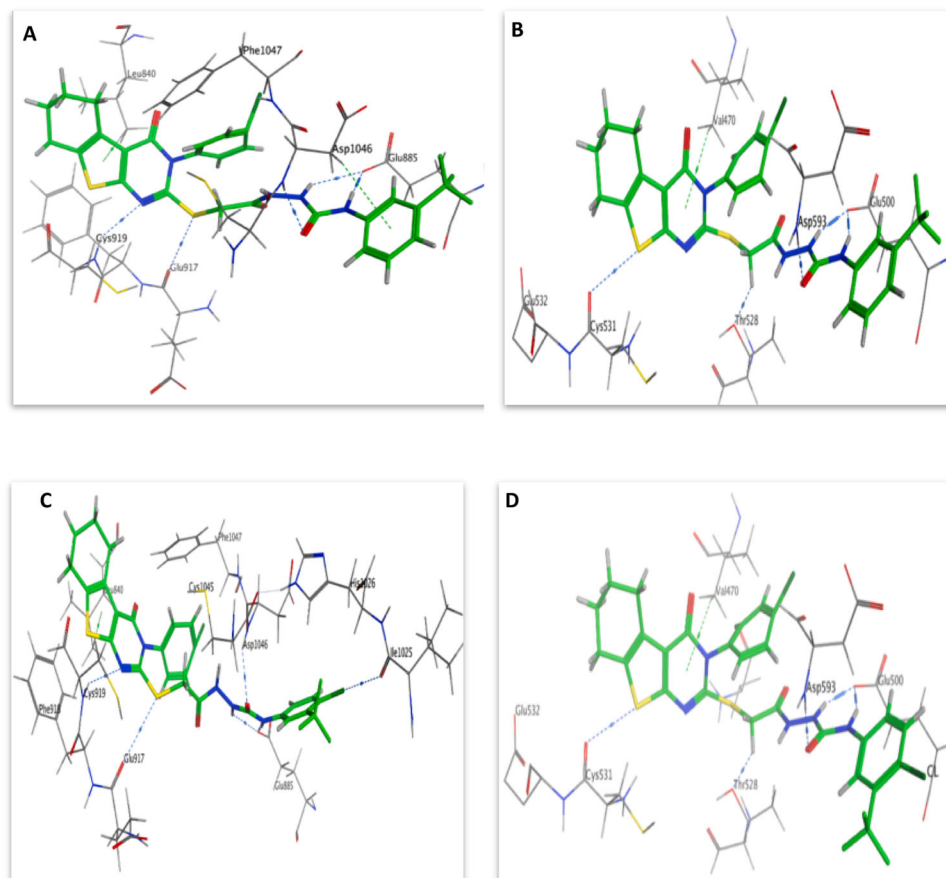


Fig. 10 SAR of thienopyrimidines as VEGFR-2/ BRAF dual inhibitors

Fig. 11 **A** Interactions of compound **40** inside VEGFR-2 binding site (PDB: 4ASD); **B** Interactions of compound **40** inside BRAF binding site (PDB: 1UWH); **C** Interactions of compound **41** inside VEGFR-2 binding site (PDB: 4ASD); **D** Interactions of compound **41** inside BRAF binding site (PDB: 1UWH)



Finally, 4-chlorophenyl cycloalkylthieno[2,3-*d*]pyrimidine formed π -H interaction with Val470 and sulfur-dipole bond with Cys531 (Fig. 11). From the mentioned data, the insertion of urea moiety in the previous compounds was crucial for VEGFR-2/BRAF activity due to the important interactions with VEGFR-2 and BRAF binding sites.

Thienopyrimidine derivatives as FMS-like tyrosine kinase-3 (FLT3) inhibitors

Early hematopoietic progenitor cells express FLT3, a type III receptor tyrosine kinase, which is essential for the survival and proliferation of hematopoietic stem cells [96, 97]. Acute myeloid leukemia (AML) is a clonal hematopoietic stem cell disease characterized by aberrant blast cell differentiation and proliferation in the bone marrow and peripheral circulation [98]. FLT3 overexpression is prevalent in AML patients, as well as other patients with FLT3 mutations [99].

Park et al. created thienopyrimidine-based analogs by modifying SPC-839, the well-known inhibitor of nuclear factor kappa-B kinase subunit beta (IKK β) to prepare derivatives **42a-e**, and then tested them in 2014 (Fig. 12) [100]. It was found that the synthesized analogs revealed no inhibitory effect against IKK β but they demonstrated

good inhibition against FLT3 kinase. Compounds **42a-e** which contain aliphatic or aromatic substituents at C-5 of thienopyrimidine revealed good FLT3 kinase inhibition with IC₅₀ ranging from 0.065 to 0.750 μ M. In addition, compound **42b** with a methyl group at C-5 showed the highest inhibitory activity on FLT3 kinase with IC₅₀ equals 0.065 μ M. Furthermore, compound **42c** with unsubstituted phenyl showed more inhibitory activity than compounds **42d** and **42e** with 3-hydroxy and 4-hydroxy phenyl substitutions. In 2016, Kim et al. developed thieno[2,3-*d*]pyrimidine derivatives to potentially act as FLT3 inhibitors for the treatment of AML (Fig. 12) [101]. Effective anti-proliferative activity against the leukemia cell line MV4-11 was shown by compounds **43a-d**, with GI₅₀ of 0.366, 0.585, 0.540, and 0.278 μ M, respectively. These compounds contain methyl group at the C-5 and cycloaminoalkoxy or elongated aminoethoxy groups in the para position of the phenyl group at C-6 which provided successful FLT3 inhibition. Therefore, compounds **43a-d** inhibited FLT3 kinase with IC₅₀ values of 3.769, 6.427, 8.026, and 2.495 nM, correspondingly. Furthermore, these compounds had improved metabolic stabilities. From the previous two studies, we can design FLT3 inhibitors upon some modification on SPC-839 structure as well as the

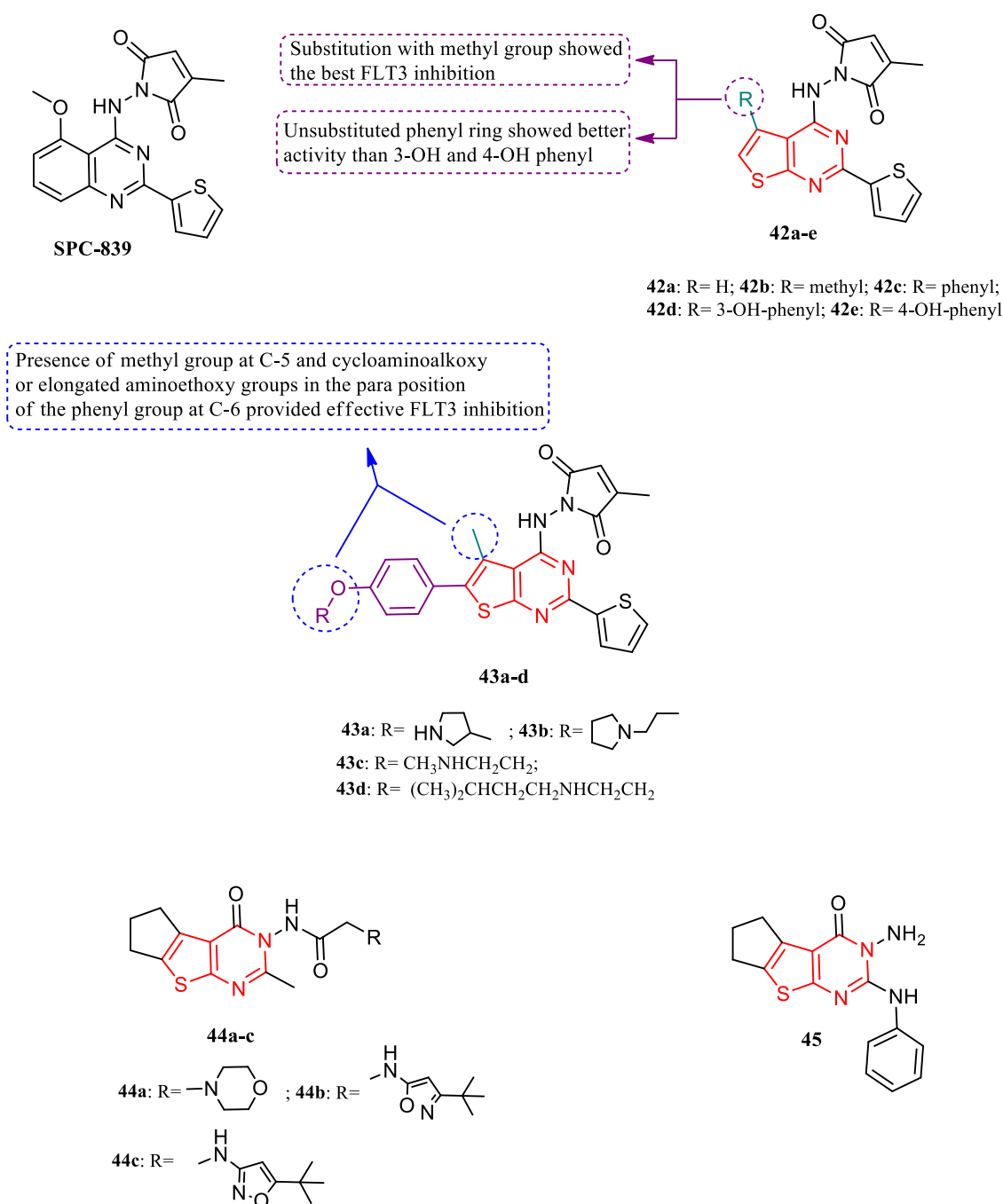


Fig. 12 SAR of thienopyrimidine derivatives as FLT3 inhibitors

presence of methyl group at C-5 of thienopyrimidine ring is valuable towards FLT3 inhibition.

Recently, in 2022 Elmongy et al. synthesized thienopyrimidine compounds that target FLT3 kinase (Fig. 12) [102]. Whereas, upon FLT3 enzyme assay, it was found that compound **44a** had the highest FLT3 inhibitory activity of the investigated compounds, with IC_{50} of $17.83 \pm 3.8 \mu\text{M}$, followed by derivatives **44b** and **45**, which had IC_{50} values of 20.4 ± 2.8 and $27.22 \pm 5.6 \mu\text{M}$, correspondingly. Moreover, compound **44c** demonstrated moderate inhibition of

FLT3 with IC_{50} of $47.64 \pm 9.3 \mu\text{M}$. On the other hand, the molecular docking study of compound **44a** inside the FLT3 active site exhibited three interactions with Leu616, Cys694, and Asp698 (Fig. 13).

Thienopyrimidine derivatives as topoisomerase II inhibitors

Topoisomerases I and II are enzymes that govern supercoiling and prevent DNA tangling, making them crucial for

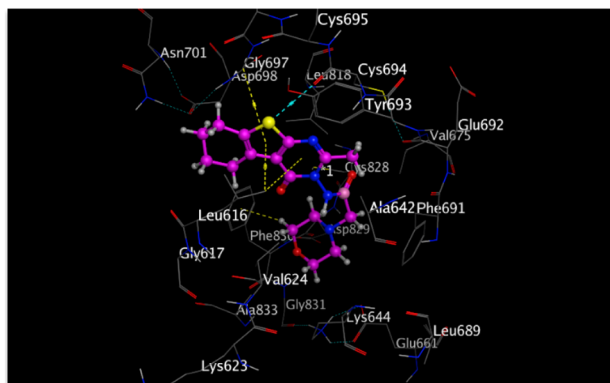


Fig. 13 Interactions of compound **44a** inside FLT3 binding site (PDB: 4XUF)

cancer cell proliferation. Topoisomerases have been identified as a key target for anticancer medicines [103]. Topoisomerase inhibitors have been developed to limit the function of topoisomerases (**I** and **II**) or to reduce their expression either or both their protein content [6, 104]. Doxorubicin, etoposide, and mitoxantrone are examples of the most well-known topoisomerase **II** inhibitors, and they serve as models for future research [105]. Abdelhaleem et al. synthesized a variety of new tetrahydrobenzothieno[2,3-*d*]pyrimidine urea derivatives (Fig. 14). Whereas, the C-4 position of thieno[2,3-*d*]pyrimidine had phenyl urea or phenyl thiourea group [106]. Consequently, compounds **46a–c** with electron-withdrawing groups showed the most powerful anticancer activity against breast cancer MCF-7 cell line with IC_{50} equals 7.10, 10.33, and 9.55 μM , respectively which were more effective than doxorubicin with IC_{50} equals 10.60 μM . Additionally, compounds **46a** and **46c** were found to be more efficient than **46b** when the electron-withdrawing group was present in the *p*-position. In addition to topoisomerase inhibition, compound **46a** inhibited many enzymes. Whereas the inhibitory activity against topoisomerase **II** with IC_{50} equals 9.29 μM and VEGFR-2 with IC_{50} equals 0.2 μM which was more effective than sorafenib. Moreover, compound **46a** significantly increased the proportion of cells in the pre-G1 and G2/M phases in comparison to control by 15.1 and 2.2 times, correspondingly, suggesting a potential role for apoptosis in compound **46a**. Furthermore, molecular modeling of **46a** showed that it could interact with essential amino acids in topoisomerase **II** binding site as NH of diaryl urea formed a hydrogen bond with Thr 744 and the carbonyl group of urea formed a hydrogen bond Tyr 734 (Fig. 15). From the previous work we can summarize the substitution of the terminal phenyl ring with electron-withdrawing group at para position is significant in anticancer activity. Recently in 2020, El-Metwally et al. synthesized and assessed thieno[2,3-*d*]pyrimidine derivatives as topoisomerase **II**

inhibitors (Fig. 14) [107]. Compounds **47–50** which contain various substituents at C-4 of thienopyrimidine showed the most potent anticancer activity against liver cancer (HepG2) and breast cancer (MCF7) cell lines with IC_{50} ranging from 4.38 to 6.71 and 3.96 to 9.19 μM , respectively. In addition, semicarbazide compound **50** significantly reduced topoisomerase **II** expression by about 60% compared to doxorubicin which reduced topoisomerase **II** expression by about 40%. Additionally, the docking of compound **50** inside the DNA binding site of topoisomerase **II** demonstrated that it interacted through the formation of two hydrogen bonds between the two amidic NH groups and AspA479. In addition, thienopyrimidine moiety created hydrophobic interactions with LysA739, ThrA783, and TyrA773. Moreover, aromatic stacking interaction was presented between LysA456 and ArgA503 and the benzene ring (Fig. 15). From the previous two studies, we assume the significant role of urea moiety in interaction with the topoisomerase **II** active site.

Thienopyrimidine derivatives as tubulin polymerization inhibitors

The cytoskeleton consists of microtubules, which play an important role in all eukaryotic cells [108]. Among their functions are cell division, mitosis, maintaining the shape of cells, motility regulation, and cell signaling [109, 110]. A successful cancer treatment approach involves creating small molecules that disrupt tubulin dynamics [111]. Tian et al. designed thieno[3,2-*d*]pyrimidine compounds as tubulin polymerization inhibitors (Fig. 16) [112]. Regarding compounds **51a–d**, it was found that the substitution of compounds **51c** (3-methyl) and **51d** (3-methoxy) with electron-donating groups enhanced the antineoplastic activity while the substitution of **51a** (3-F) and **51b** (3-Br) with electron-withdrawing groups decreased the antineoplastic activity. Therefore, compound **51c** revealed effective tubulin polymerization inhibition with IC_{50} of $4.1 \pm 0.1 \mu\text{M}$. Additionally, a cell cycle study of **51c** revealed that it caused G2/M arrest in Hela cells. According to molecular modeling of compound **51c** in tubulin, the 3-methyl group on the phenyl ring was tightly positioned within a sub-pocket shaped by Val181 (α -monomer) and Val315 (β -monomer) (Fig. 17). From this study, we can conclude that the presence of methyl group in position 3 of the phenyl ring significantly increased the antiproliferative action and it was important for tubulin polymerization inhibition due to its interaction with the active site of tubulin. In 2018, Yang et al. synthesized thienopyrimidine derivatives having dithiocarbamate moiety at C2 (Fig. 16) [113]. The results showed that compounds substituted with strong electron-withdrawing groups at position 4 of terminal phenyl ring as **52b** (CN) and **52c** (NO_2) were more

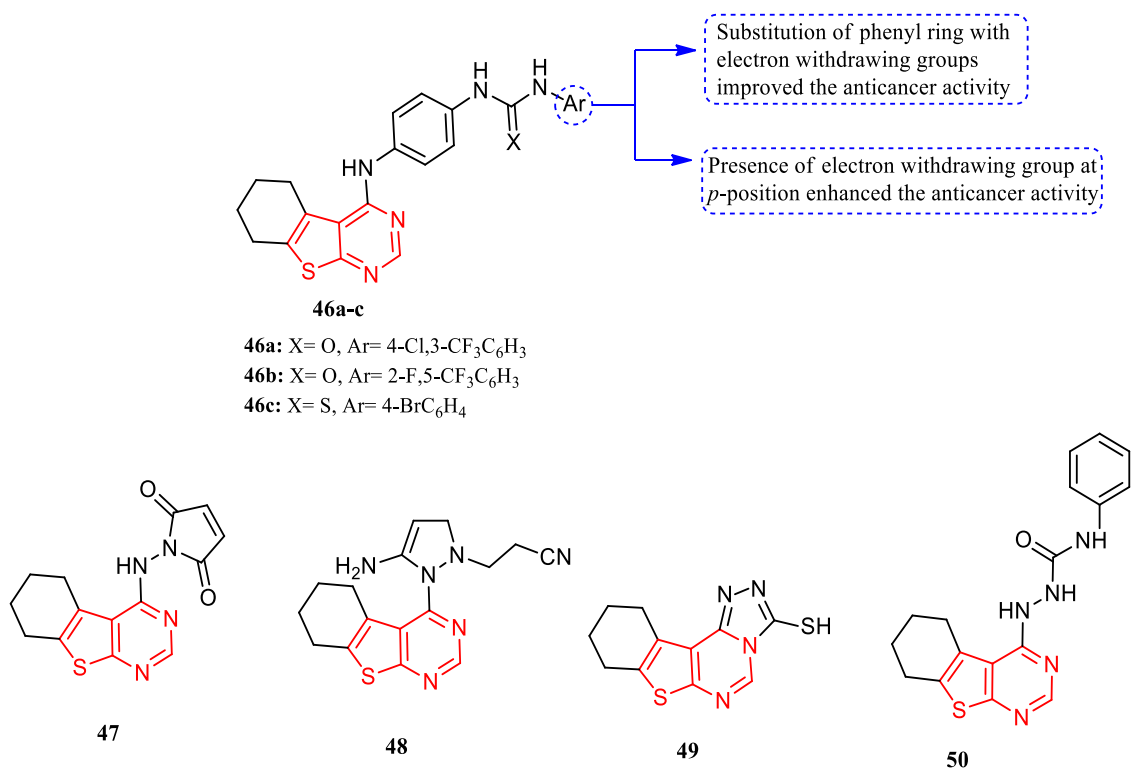
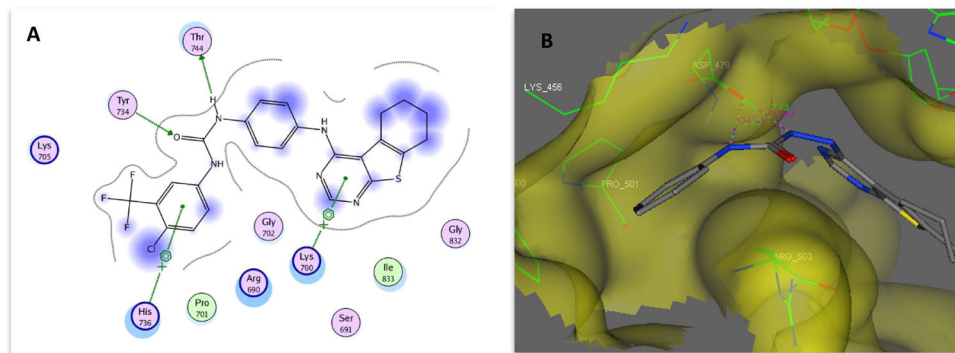


Fig. 14 SAR of thienopyrimidines as topoisomerase II inhibitors

Fig. 15 **A** Interactions of compound **46a** inside the DNA binding site of topoisomerase II (PDB: 1ZXM); **B** Interactions of compound **50** inside the DNA binding site of topoisomerase II (PDB: 3qx3)



powerful than compound **52a** with unsubstituted phenyl ring. On the other side, compounds substituted with electron-donating groups at position 4 of the phenyl ring as **52d** (methyl) and **52e** (methoxy) demonstrated comparable anticancer activity with **52a**. Hence, compound **52b** presented the highest antineoplastic activity against A549 cell line with IC₅₀ of 4.87 μM. As a result of **52b**, caused cell cycle arrest at the G2/M phase and the spindle assembly checkpoint (SAC) is activated. Furthermore, compound **52b** showed tubulin polymerization inhibition in a dose-dependent manner. From this study, we can highlight the importance of the substitution of phenyl ring with a strong electron-withdrawing group to achieve the maximum anticancer activity.

Thienopyrimidine derivatives as histone deacetylase inhibitors

It has been well-documented that HDACs play an important role in epigenetic regulation [114]. Histone acetylation is regulated by two enzymes: HDACs and histone acetyltransferases (HATs) [115]. Normally, HATs and HDACs are balanced in cells. Alternatively, cancer cells have an imbalance between histone acetylation and deacetylation due to overexpression of HDACs or suppression of HATs, resulting in oncogene activation and tumor progression [116]. Therefore, Developing novel anticancer agents through the inhibition of HDACs has proven successful [117]. In 2017, Wang et al. designed thienopyrimidine

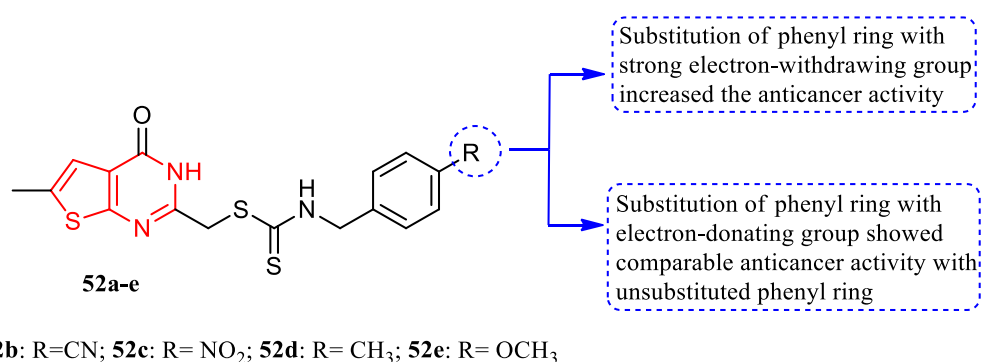
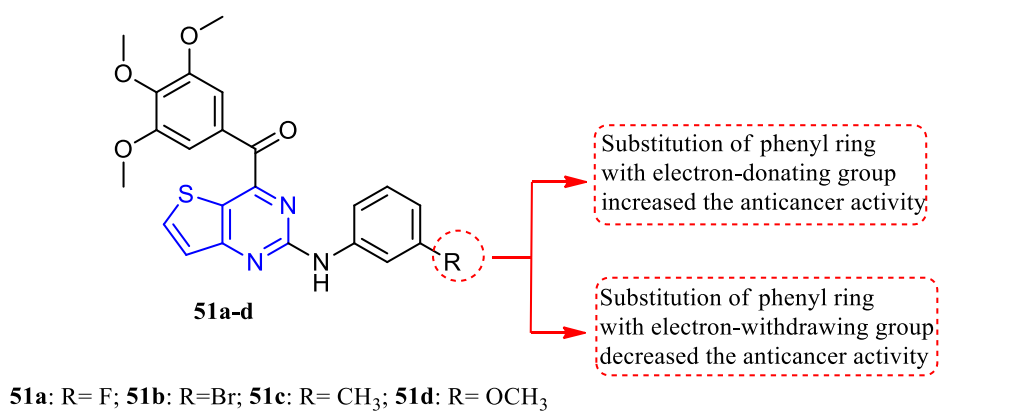


Fig. 16 SAR of thienopyrimidine derivatives as tubulin polymerization inhibitors

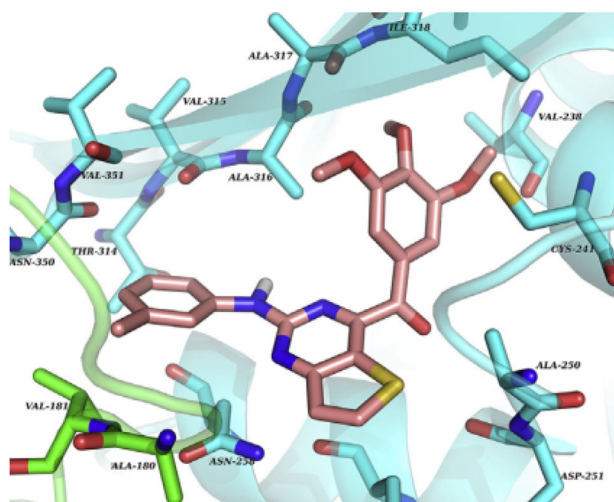


Fig. 17 Interactions of compound **51c** inside tubulin binding site (PDB: 5LY)

derivatives containing hydroxamic acid as HDAC inhibitors (Fig. 18) [118]. The design of the synthesized compounds (**53a-c**) is based on the presence of hydroxamic acid which is an essential functional group for HDACs inhibition and

act as zinc binding group (ZBC) at the HDACs active site. The results of tested compounds demonstrated that the introduction of a bulky group at position 3 or 4 of phenyl ring as in **53b** (3-tertbutyl) and **53c** (4-isopropyl) decreased HDACs inhibition. Finally, compound **53a** (3-ethynyl) was observed to be an effective inhibitor of HDAC1, HDAC3, and HDAC6 with IC₅₀ values of 29.81 nM, 24.71 nM, and 21.29 nM, respectively. Mohamed et al. synthesized thienopyrimidine compounds which were evaluated as HDAC inhibitors (Fig. 18) [119]. Consequently, compounds containing hydroxamic acid as a zinc-binding group (ZBC) either with an aliphatic or aromatic linker (**54a** and **55**) exhibited high inhibitory activity against HDAC while replacement of hydroxamic acid moiety with hydrazide (**54b**) or 2-aminoanilide (**54c**) groups reduced HDAC inhibition. Moreover, it was observed that compounds with aliphatic linkers were more effective against HDAC than compounds with aromatic linkers as illustrated in. Therefore, compound **54a** revealed the most effective HDACs inhibition with IC₅₀ against HDAC1, HDAC2, HDAC6, and HDAC8 equals 0.028 μM, 0.078 μM, 0.471 μM, and 1.903 μM, respectively. Additionally, a molecular docking study of compound **54a** with HDAC2 active site exhibited

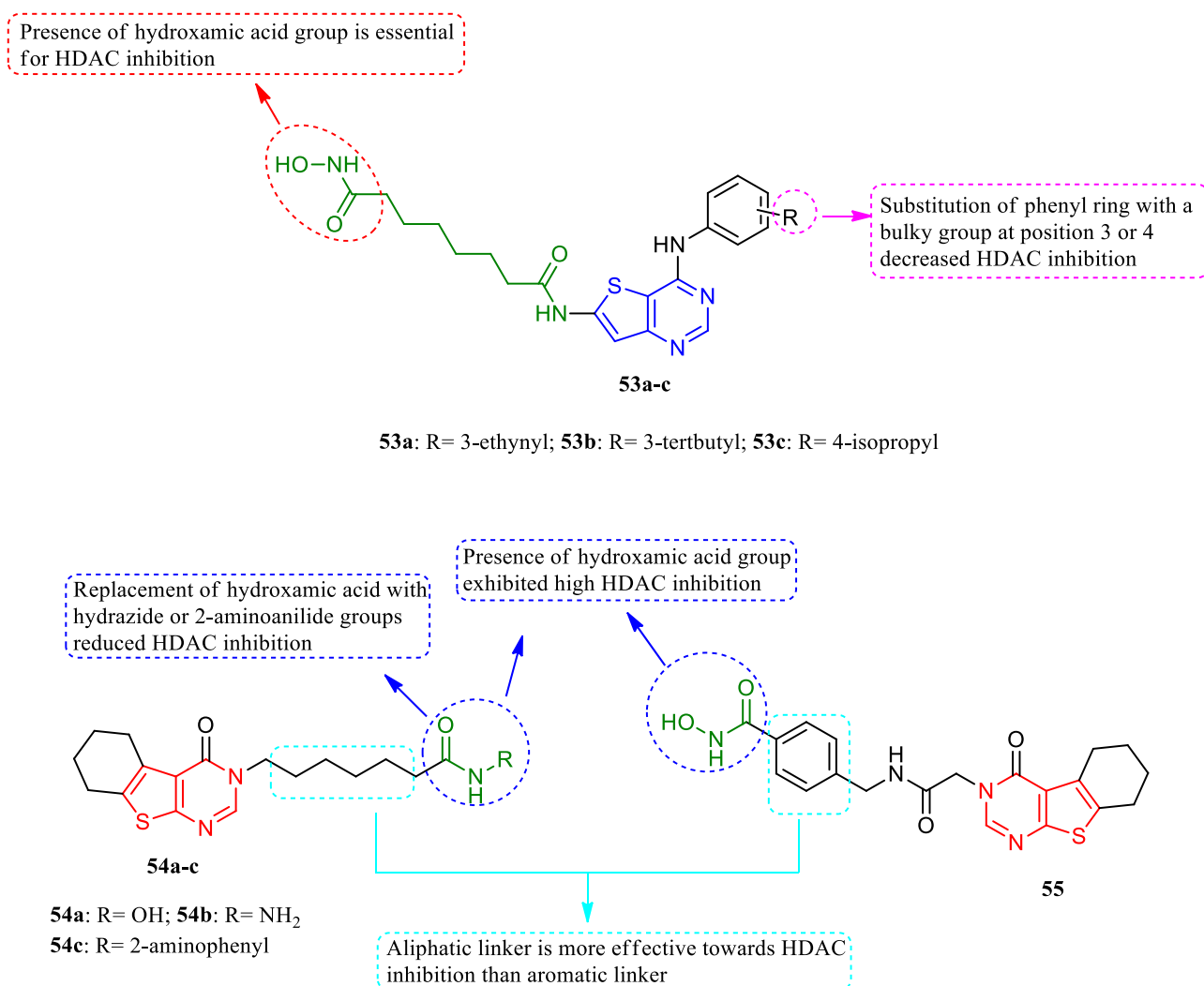


Fig. 18 SAR of thienopyrimidine derivatives as HDAC inhibitors

that the hydroxamic acid group (ZBG) formed hydrogen bonds with His145, His146, and Tyr308. Besides, it showed metallic bonds with Zn ion and hydrophobic interaction with His33 (Fig. 19). From the previous research, we can assume that the presence of a hydroxamic acid group with an aliphatic linker achieved significant HDAC inhibition and it is a favorable feature for designing HDAC inhibitors.

Conclusion

Thienopyrimidine derivatives perform a significant role in the production of drugs that have different pharmacological activities particularly, anticancer activity. Thienopyrimidines act as anticancer agents through diverse enzyme inhibition as (EGFR, VEGFR-2, BRAF, etc.). So in this review, the most recent publications on thienopyrimidine scaffold synthesis and anticancer evaluation have been reviewed. The current review can help scientists and

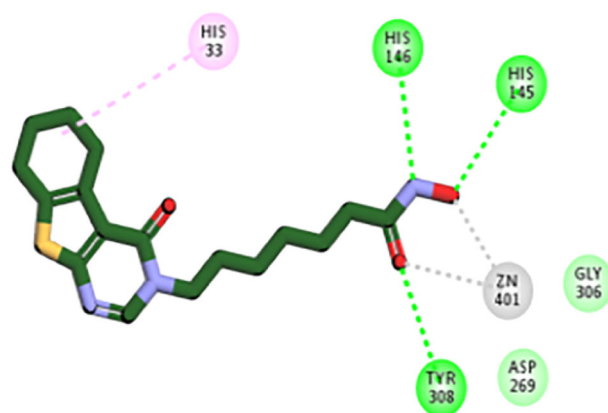


Fig. 19 Interactions of compound 54a inside HDAC2 binding site (PDB: 4LXZ)

researchers from around the world select precisely the goals for the future development of powerful lead compounds as antineoplastic medicines.

Acknowledgements The authors are thankful to the Faculty of Pharmacy, Cairo University.

Funding Open access funding provided by The Science, Technology & Innovation Funding Authority (STDF) in cooperation with The Egyptian Knowledge Bank (EKB).

Compliance with ethical standards

Conflict of interest The authors declare no competing interests.

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