



REVIEW ARTICLE

EXPLORING UTILITY OF PYRAZINE-BASED HETEROCYCLIC COMPOUNDS IN ANTICANCER DRUG DEVELOPMENT

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Cancer is witnessing a rising global incidence, affecting a predominant proportion of the populace. In response, ongoing endeavors are underway to develop novel anticancer pharmaceuticals, with their safety profiles currently undergoing evaluation. Among these, pyrazine-based medications constitute a significant contribution, representing a pivotal pharmacophore prevalent in both synthetic and natural heterocyclic compounds. Characterized by a six-membered aromatic heterocycle with two nitrogen atoms, pyrazine exhibits versatile therapeutic applications in drug development, offering myriad prospects for the refinement of future anticancer agents through targeted interactions with pivotal receptors. Pyrazine compounds, through their inherent structural attributes, have demonstrated efficacy in inhibiting enzymes, receptors, and various other targets implicated in cancer pathogenesis. Contemporary research endeavors are centered on the synthesis of novel pyrazine derivatives tailored for cancer treatment, often in combination with other molecular moieties. Consequently, this review expounds upon the recent therapeutic advancements in pyrazine-based drugs, providing insights into their synthetic pathways, marketed drugs elucidating their primary targets, and a compendium of recently patented and published research papers. This collective information serves as a valuable resource for scientists engaged in the formulation of efficacious medications endowed with the requisite pharmacological activity.

Key words: Heterocyclic compounds, Cancer, Pyrazine-based drugs, Anticancer pharmaceuticals.

INTRODUCTION

Pyrazine is an azaheterocycle that is aromatic and contains two nitrogen atoms. Specifically, pyrazine is a diazine and is isomeric with pyrimidine and pyridazine. Pyrazine is an electron-deficient molecule because of the presence of two electronegative N-atoms which withdraws electron density from the aromatic ring. Pyrazine derivatives are well-known in natural products and are reported to have medicinal and biological significance. In the laboratory, it was first synthesized in 1876 [1]. Pyrazine is an important pharmacophore since it constitutes the integral scaffold in several types

of drugs. Although literature is enriched with reports of cytotoxic potential of heterocyclic and other aromatic compounds [2, 3], but due to the wide bio-spectrum of pyrazine derivatives, this motif is present in anti-inflammatory, anticancer, antidiabetic, and diuretic drugs [4].

Furthermore, pyrazine derivatives are also reported to possess high antimicrobial activities [5]. In addition to the importance of pyrazine derivatives in the pharmaceutical sector, these molecules have immense commercial importance in flavor [6], fragrance [7], and food industries [8]. Pyrazine, a naturally occurring

compound in various living organisms, primarily serves as a flavor enhancer in raw foods. Beyond its natural occurrence, industries produce pyrazine and its derivatives in fragrances, flavors, and pharmaceuticals. This review delves into the historical evolution of pyrazine, from its discovery to recent synthetic methodologies, notably focusing on the synthesis of pyrazinium. Six primary synthetic approaches are explored: condensation reactions, ring closure, metal catalysis, green reactions, the Maillard reaction, and acid-catalyzed N-substitution. The initial five methods primarily target substitutions at 2nd, 3rd, 5th, and 6th positions in the pyrazine ring, while the last method specifically focuses on positions 1 and 4 in the pyrazine structure [9]. Pyrazine, a type of six-membered nitrogen-based heterocycle, is well-known for its application in rational drug development [10]. Its ring ligands have been intensively explored, and their p-donor properties are promising [11]. It is an aromatic heterocyclic compound having 1,4-substituted benzene nitrogen atoms used as tastes, fragrances, medicines, and chemical raw materials [12]. Many natural pyrazines have been discovered over the last 70 years [13, 14]. It has received much attention because of the diazine rings, a beautiful class of molecular structures found in both natural and synthesized molecules [15]. The bioactive properties of target molecules are significantly altered by structural modifications in the pyrazine moiety. As a result, ongoing attempts have been made to synthesize a wide class of pyrazine derivatives that may be effective in the battle against cancer cells [16]. Even pyrazine-containing compounds have an important role in everyday life, having applications in materials science [17] and pharmaceutical chemistry [18-23] (**Figure 1**).

General synthesis of Pyrazine

Since pyrazine is a therapeutically important moiety, its contribution is increasing widely in drug discovery. It can be synthesized in various ways; some of the reported synthesis schemes are mentioned below (**Schemes 1-4**).

Anti-tumor activity of pyrazine derivatives

BRAF and tubulin inhibitors

The B-Raf protein is produced by the BRAF gene in humans; the protein is called serine/threonine-protein kinase B-Raf. The B homologs of the murine sarcoma viral oncogenes B-Raf and v-Raf are proto-oncogenes [30, 31]. The B-Raf protein is in charge of delivering signals that

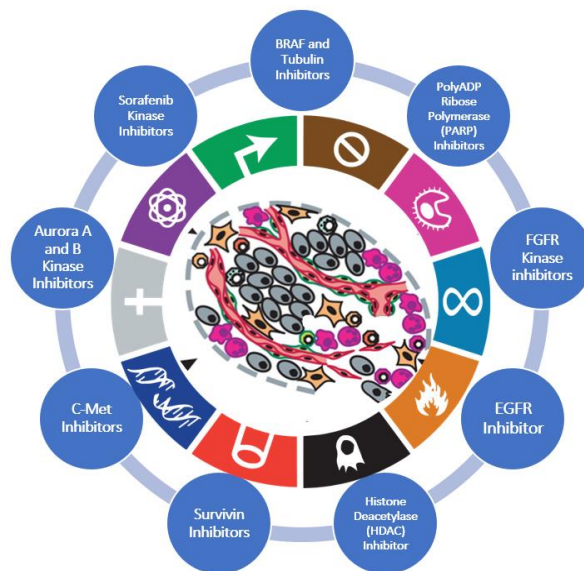


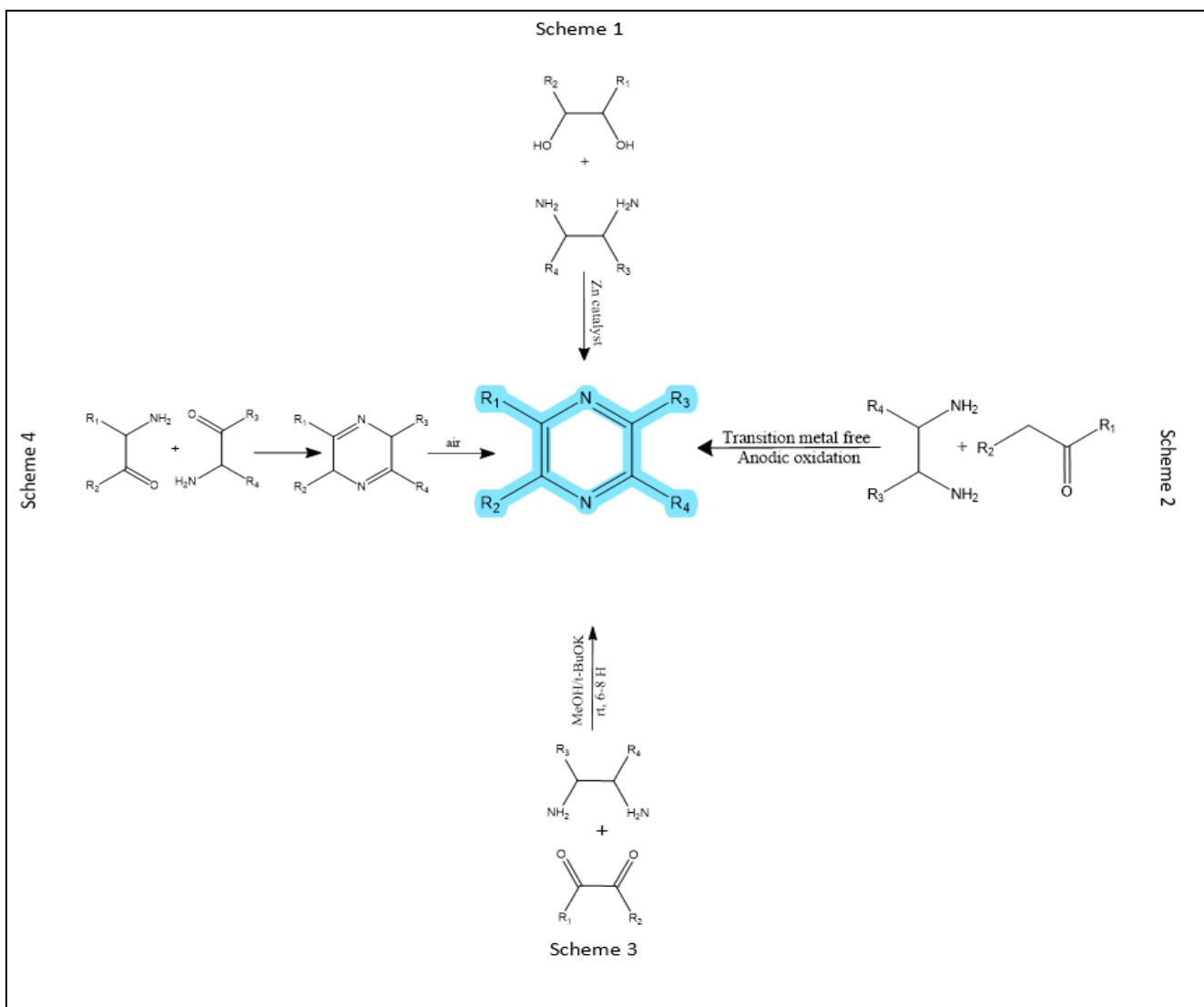
Fig. 1. Diverse activities of pyrazine derivatives

govern cell growth inside cells. It was discovered to be mutated in some human malignancies in 2002 [32]. Tubulin is a component in the cytoskeleton of the eukaryotic cells that plays a role in intracellular movement and transport, as well as cell signaling, mitosis, polarity, and cell shape maintenance [33]. They are also involved in the process of cell division by assisting in the attachment and movement of chromosomes throughout numerous phases of mitosis. As a result, microtubule dynamics is a key target for anti-tumor therapy development [34].

A series of compounds containing the pyrazino [1, 2-*a*] indol-1(2*H*)-one pharmacophore and evaluated their anticancer activity against a panel of cancer cell lines [35]. Compounds 1c and 1d were the most effective with inhibitory IC_{50} values of (1c, 2.4 ± 1.2 and 1.0 ± 0.2) and (1d, 1.7 ± 0.5 and 0.1 ± 0.07 , 0.6 ± 0.2 μ M) against EGFR and BRAF, respectively. Similarly, the tubulin effect of 1c and 1d compounds were satisfactory at 930 ± 242 and 650 ± 227 (arbitrary units), respectively. These most potent compounds have methyl (1d) and hydrogen (1c) substitutions at the pyrrole ring of the indole moiety and chloro-substitutions at the aromatic ring, making them the best inhibitory and antiproliferative agents. Furthermore, the indole carboxylic amide moiety is crucial for the ROS inhibitory effect, and only these two compounds, 1c and 1d, had a substantial impact on tubulin assembly (**Figure 2**).

PolyADP ribose polymerase (PARP) inhibitors

PARP inhibitors are medications that stop the enzyme poly ADP-ribose polymerase (PARP)

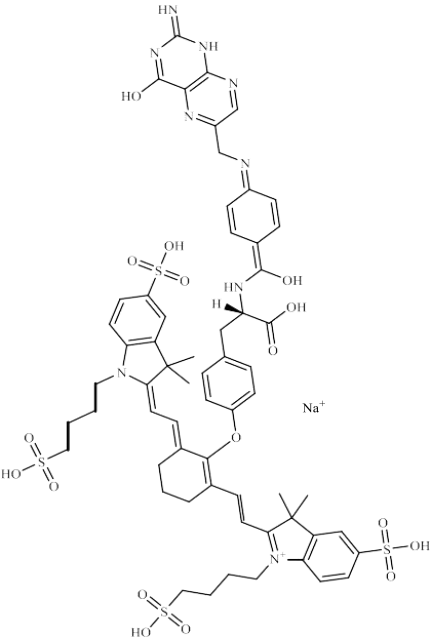
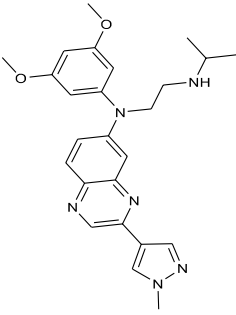
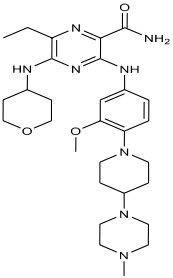
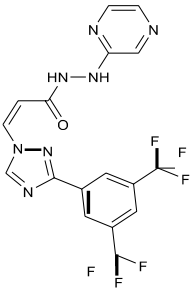


Sch. 1. Formation of pyrazine in the presence of zinc was reported by Sato (1978) [24]; **Sch. 2.** Synthesis of pyrazine *via* electrochemical dehydrogenative [4+2] annulation between ketones and diamines [25]; **Sch. 3.** The green approach in producing pyrazine was reported by Ghosh and Mandal (2012) [26, 27]; **Sch. 4.** Synthesis via interaction with ammonia, an aminoketone is formed, which is subsequently condensed and oxidized to pyrazine [28]

Further, a list of pyrazine-based drugs approved by the US FDA for cancer treatment with their primary purpose are listed in **Table 1**.

Table 1. A list of drugs approved by the US FDA for cancer treatment with their primary purpose [29]

S. No.	Drug	FDA approval	Primary target	Company and brand	Structure
1.	Acalabrutinib	Nov 21, 2019	Binds cysteine residue in the BTK	CALQUENCE, AstraZeneca	

2.	Pafolacianine	Nov 29, 2021	Folate receptor	CYTALUX, On Target Laboratories, LLC	
3.	Erdafitinib	Apr 12, 2019	FGFR phosphorylation and signaling	BALVERSA, Janssen Pharmaceutical Companies	
4.	Gilteritinib	Nov 28, 2018	Tyrosine kinases	XOSPATA, Astellas Pharma US Inc	
5.	Selinexor	Jun 22, 2020	Nuclear export of tumor suppressor proteins (tsps)	XPOVIO, Karyopharm Therapeutics	

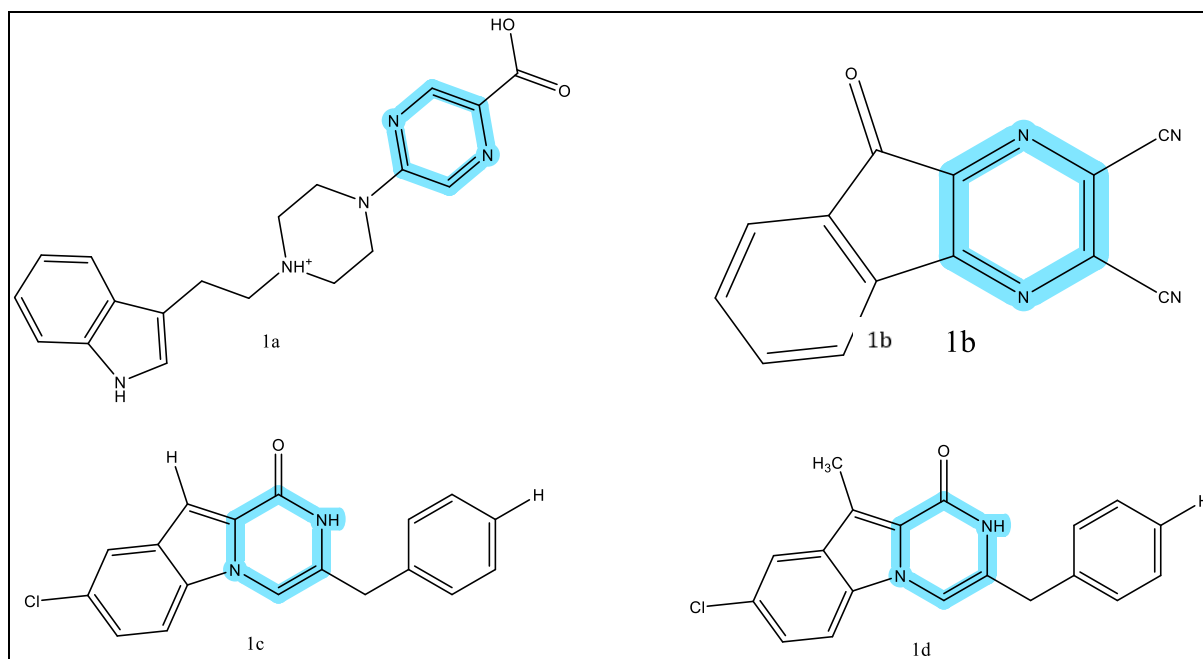


Fig. 2. Structure of compounds 1a-d

from working. They're being created for a variety of applications, including treatment of hereditary malignancies [36]. PARP (PARP1, PARP2, etc.) is more important in cancer cells as compared to normal cells, making it a very important target in cancer therapy [37, 38]. As indicated mostly by olaparib coupled to standard treatment, in women who have recurrent platinum-sensitive ovarian cancer, PARP inhibitors appear to improve progression-free survival [39].

The synthesis of 3,4- dihydropyrrolo[1,2-*a*] pyrazine derivatives via the [4+1+1] annulation

approach is reported with evaluation of their anti-cancer properties [40]. The cytotoxic effect of compound 2b was the highest, with IC_{50} values of $1.18 \pm 0.05 \mu\text{M}$ and $1.95 \pm 0.04 \mu\text{M}$ against PC-3 and MCF-7 cell lines, respectively. Also, it significantly decreased cell migration in a dose-dependent manner and caused apoptosis in PC-3 and MCF-7 cells by activating caspase-3 and cleaving PARP. Furthermore, a phenacyl group is present at the nitrogen of the pyrazine ring in compound 2b, due to which it more potently decreases cell viability than other derivatives (**Figure 3**).

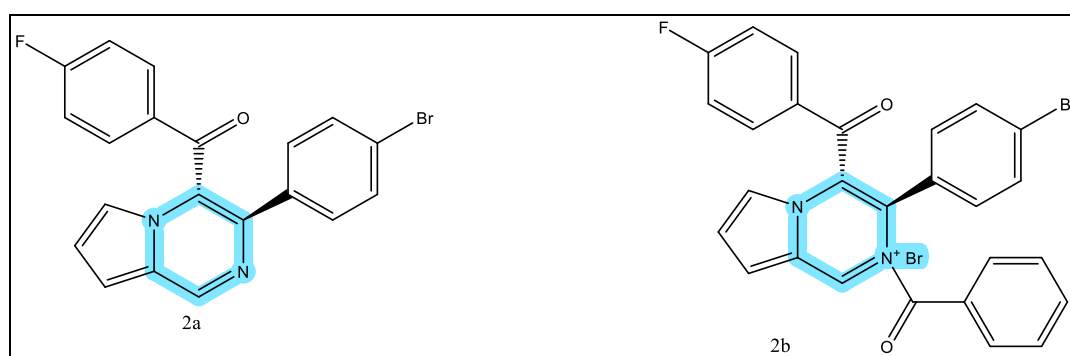


Fig. 3. Structure of compounds 2a and 2b

FGFR kinase inhibitors

Recently, the signaling pathways initiated by FGFs have been discovered to be very critical in the development and progression of various malignancies. The fibroblast growth factor receptors (FGFR) bind to members of the FGF family of proteins, as their name implies [41-44].

Pathological disorders involve some of these receptors. The small molecule inhibitors are divided into non-selective and selective FGFR inhibitors. They all target the ATP-binding site in the kinase domain [45]. A class of 5H-pyrrolo[2,3-*b*] pyrazine-based derivatives as strong FGFR inhibitors, the best of those had

sub-nanomolar enzymatic activity [46]. Among all, chemical 3d was found to be effective against tumors with fibroblast growth factor receptors (FGFR) genetic alterations, with IC_{50} values of 0.6 ± 0.0 and 0.5 ± 0.3 nmol/l against FGFR1 and KG1, respectively. Furthermore, compound 3d possesses a dimethoxybenzene ring that,

by extending from methyl to ethyl or cyclist groups, drastically reduces the activity of inhibition. The fluorine group in the dimethoxybenzene ring slightly increased the IC_{50} activity to 0.4 nmol/l. These potentials of compound 3d make it more conducive to further development (**Figure 4**).

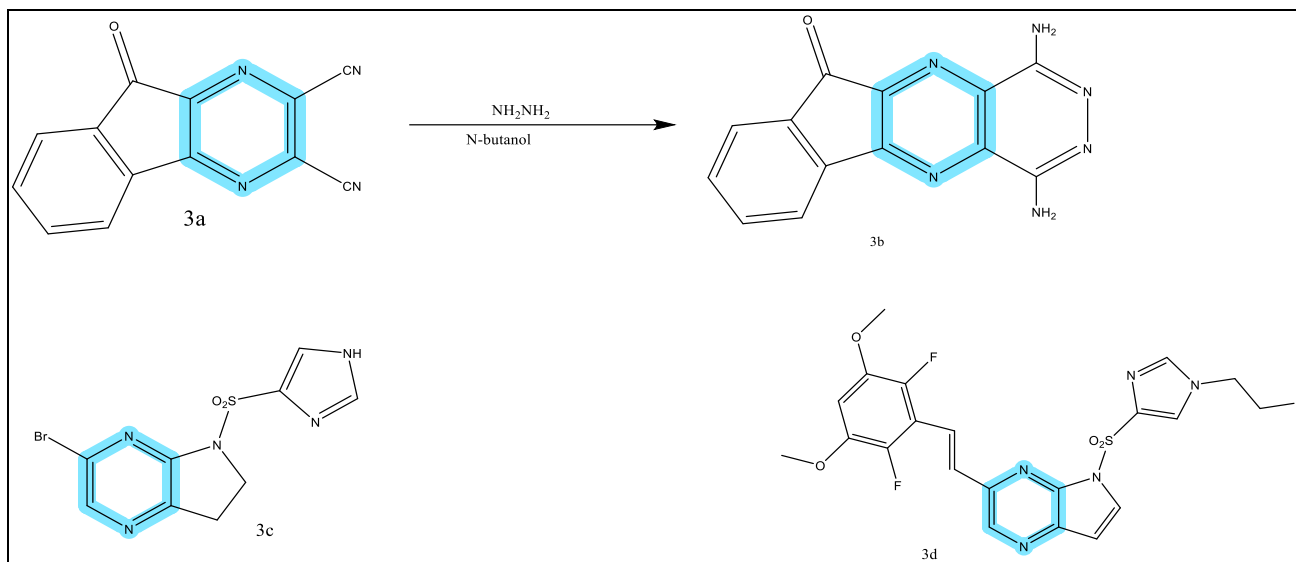


Fig. 4. Structure of compounds 3a-d

Numerous effective FGFR kinase inhibitors, including a series of 5*H*-pyrrole [2, 3-*b*] pyrazine derivatives that were rationally designed, synthesized, and biologically evaluated [47]. Compound 3e was a promising lead for further development due to its excellent selectivity and favorable metabolic characteristics, with an $IC_{50} = 0.6 \pm 0.2$ nM against FGFR1. Furthermore, compound 3e has an unsubstituted pyrazole ring, which showed much higher activity than others when isopropyl, carbonyl, or any other group was introduced to the pyrazole ring. Its activity decreased drastically against FGFR1. In addition, the docking study also showed that compound 3e has excellent binding affinity to the crystal structure of FGFR1 due to its imidazole, pyrazole, & pyrazine rings (**Figure 5**).

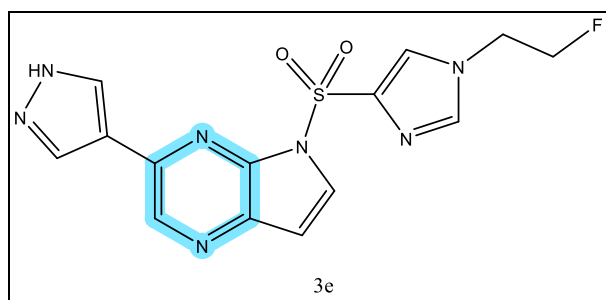


Fig. 5. Structure of compound 3e

EGFR Inhibitor

The epidermal-growth-factor receptor (EGFR/HER1/ ErbB-1) is a transmembrane protein that serves as a receptor for extracellular proteins belonging to the EGF family [48]. The ErbB family of receptors comprises four closely associated tyrosine kinase receptors: HER2/neu (ErbB-2), ErbB-1, Her 4 (ErbB-4), and Her 3 (ErbB-3). Tumors can be caused by mutations that affect EGFR expression or function in a range of cancers [49] reported the synthesis of fused quinoxaline and pyrazine derivatives, which were cytotoxicity tested against MCF-7 and A549 cell lines was reported [50]. A compound named 2b, obtained by reacting compound 2a with hydrazine hydrate in n-butanol, was found to be the most active compound, with IC_{50} values of 4.3 and 5.4 μ M against A549 and MCF-7, respectively. A study on molecular docking also revealed that compound 2b exhibited a good binding affinity towards the EGFR receptor binding site (PDB ID: 1M17).

Histone deacetylase (HDAC) inhibitor

HDACs catalyze the removal of acetyl groups from histones, which leads to heterochromatin (condensed chromatin) and gene transcription

inhibition [51]. Based on sequence homology and domain organization, human HDACs are split into four groups. Class-I (HDACs 1, 2, 3, and 8) zinc-dependent deacetylases, class-II (HDACs 4, 5, 6, 7, 9, and 10) zinc-dependent deacetylases, and class-IV (HDAC11) NAD⁺-dependent deacetylases (sirtuins SIRT1-7) [52]. HDACs 1, 2, 3, and 8 are predominantly present in the nucleus and are involved in cell proliferation cell cycle progression, maintenance, and also in the formation of cancer cell phenotypes [53]. HDAC inhibitors (HDACi) have been created and explored as possible anticancer therapies in recent years since HDACs are critical for tumor formation and progression [54].

The synthesis of a family of class I selective

HDAC inhibitors (HDACi) that have a zinc-binding group attached to a central (piperazin-1-yl) pyrazine scaffold [55]. Compounds 5a, 5b and 5c were shown to be the most active compounds against HDAC 1, 2, and 3, with IC₅₀ values of (0.13, 0.28, 0.31 μM), (0.26, 2.47, 0 μM), and (0.07, 0.26, 6.1 μM), respectively. Furthermore, evaluation of these compounds showed that 1c is superior; the inhibitory action against HDAC is attributed to its 3-indolyl ring, and its activity is decreased by the replacement of the methylene that connects the indole and piperazine groups with ethylene. The docking study also revealed a good interaction of compound 5a with the crystal structure of HDAC (PDB IDs: 4BKX, 4LY1, and 4A69) (Figure 6).

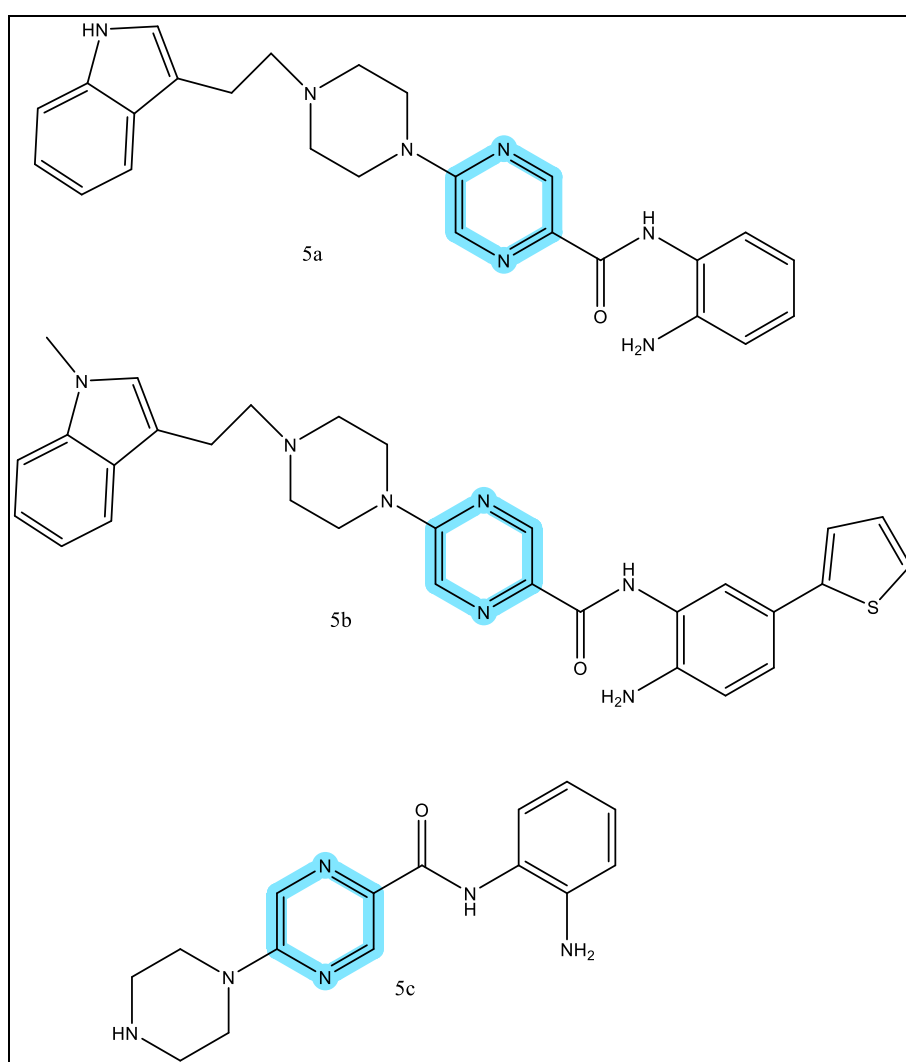


Fig. 6. Structure of compound 5a-c

Survivin inhibitors

Survivin is a protein that is also known as BIRC5 (baculoviral inhibitor of apoptosis repeat-containing 5), which is represented by the BIRC5 gene in humans [56, 57]. It belongs to the

inhibitor of apoptosis (IAP) family, and its function has been conserved throughout evolution, as homologs of the protein have been found in both vertebrates and invertebrates [58]. In most human Tumors and foetal tissue,

the survivin protein is widely expressed but is completely absent in terminally differentiated cells [59]. A tetracyclic aromatic derivative (compound 6b) using a modified version of a previously published compound 11a and assessed its anti-tumor efficacy [60]. It has greatly enhanced potency, with IC₅₀ values of 0.16 and 0.17 M against PC-3 and C4-2 cells, and causes prostate cancer cells to spontaneously

apoptosis. It also more potently inhibits survivin dimerization and promotes survivin breakdown. The 1,2,5-oxadiazole moiety does not appear to be necessary as it interacts moderately with survivin; therefore, its replacement or removal changes the effectiveness. Furthermore, the docking simulation with a human survivin (PDB ID: 1F3H) similarly demonstrated the 6b's increased potency (Figure 7).

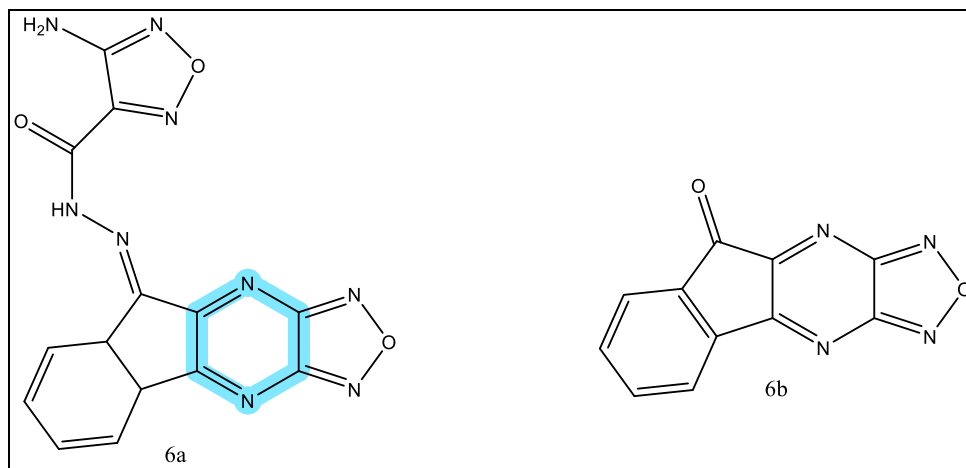


Fig. 7. Structure of compounds 6a and 6b

C-Met inhibitors

Tumor activity against the cell lines - Hela (oldest and commonly used human cell line), A549, and MCF-7 with IC₅₀ values of 2.85±0.74, 0.83±0.07, and 0.15±0.08 μM, respectively. c-Met-kinase inhibition was also superior at the nanomolar level (IC₅₀=48 nM). Additionally, compound 7a showed a greater antitumor

activity than other compounds due to the presence of a bigger electron-withdrawing group (trifluoromethyl group) on the phenyl ring. Furthermore, the docking investigation demonstrated its potential for binding to the crystal structure of c-Met (PDB ID: 3LQ8). The findings suggested that it could be a potential inhibitor (Figure 8).

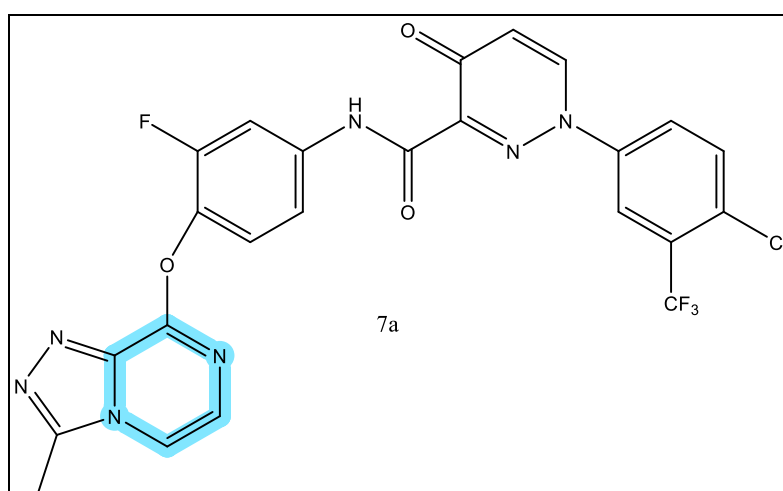


Fig. 8. Structure of compound 7a

In healthy people, the (HGFR), commonly known as C-Met, is involved in invasive growth during embryonic development, wound healing, and tissue repair [61-63]. Tumor development and

metastasis are mediated by abnormal C-Met signaling. As a result, a variety of strategies have been used to reduce the aberrant c-Met axis as an anticancer therapy [64-66]. Small molecule

inhibitors have gotten a lot of attention among them [67, 68].

The *c*-Met kinase inhibitory and antiproliferative properties of several 5-methylpyrazolo[1,5-*a*]pyrimidine derivatives were reported [69]. The majority of the drugs inhibited *c*-Met kinase with remarkable potency and selectivity against the cancer cell lines. The compounds with the most active antiproliferative effect were compounds 7c and 7b, which had IC_{50} values of 7c (26.83 ± 2.41 and 73.33 ± 1.75) and 7b (93.73 ± 4.24 and 20.20 ± 2.04 μ M) against MDA-MB-231 and A549, respectively. Accordingly, compound 7c showed to be the most effective *c*-Met inhibitor, with an IC_{50} value of 5.170.48 nM, while compound 7b showed to be the second

most effective inhibitor, with an IC_{50} value of 5.620.78 nM. Furthermore, compound 7c includes a pyrazine moiety, which increases its potency. Similarly, compound 7b has 4-fluorophenyl, which likewise shows good cytotoxicity toward all cell types. These findings revealed that the addition of fluorophenyl and pyrazine rings was advantageous for broad-spectrum anticancer activity, even if other benzene ring substitutions, such as methyl, chloro, metamethoxy, and 3-chloro-4-methoxy, decreased cytotoxicity. Additionally, docking analysis revealed that substituting pyrazine might enhance the ability to inhibit *c*-Met kinase and the antiproliferative potential (PDB IDs: 3U6I and 4KNB) (Figure 9).

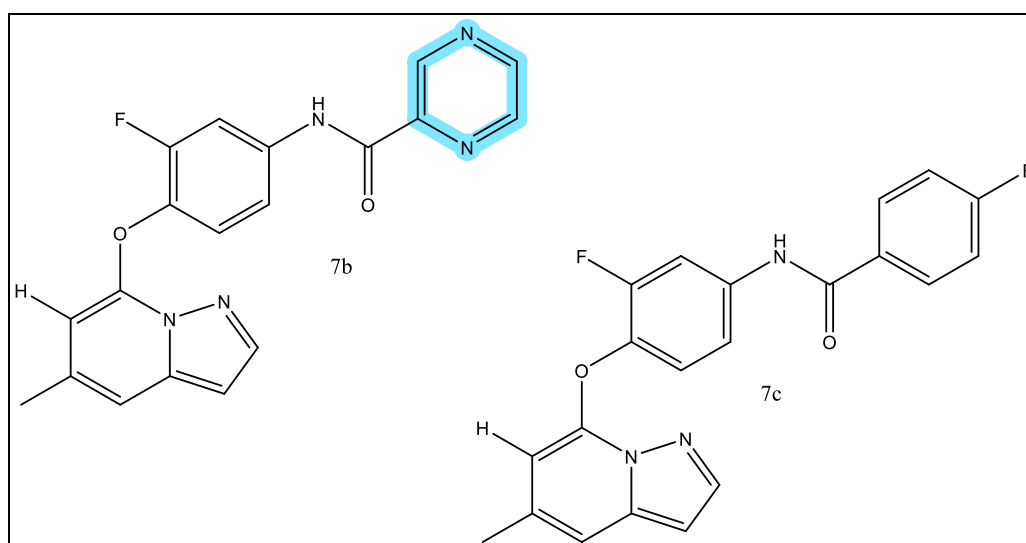


Fig. 9. Structure of compound 7b, 7c

A series of [1,2,4]-triazolo [4,3-*a*]-pyrazine compounds containing the 4-oxo-pyridazinone moiety and tested them against cancer cell lines as well as the *c*-Met kinase [70]. The most potent chemical, 10e, showed excellent anti-tumor activity against the cell lines Hela, A549, and MCF-7 with IC_{50} values of 2.85 ± 0.74 , 0.83 ± 0.07 , and 0.15 ± 0.08 μ M, respectively. *c*-Met-kinase inhibition was also superior at the nanomolar level ($IC_{50}=48$ nM). Additionally, compound 10e showed greater antitumor activity than other compounds due to the presence of a bigger electron-withdrawing group (trifluoromethyl group) on the phenyl ring. Furthermore, the docking investigation demonstrated its potential for binding to the crystal structure of *c*-Met. This suggested that it could be a potential inhibitor.

Aurora A and B kinase inhibitors

The aurora kinase family is a serine / threonine

kinase subfamily that controls centrosome maturation, mitotic spindle formation [71], cytokinesis, and chromosomal segregation during mitosis [72]. Aurora A inhibition causes mitotic latency and the production of monopolar spindles, which leads to cell death [73], while aurora B is a transcription factor that controls chromatin remodeling, kineto- chore-spindle attachment, and cytokinesis [74]. Inhibition of Aurora B causes endoreduplication and cytokinesis to be arrested, resulting in apoptosis. When both aurora A and B are blocked, the inhibition phenotype of aurora B dominates [75]. The antiproliferation of 3,5-disubstituted-2-aminopyrazines as aurora kinase inhibitors was reported [76]. Among all, compound 8a had the highest antiproliferative activity against HepG2, LoVo, HeLa, and U38 cells, with IC_{50} values of 7.30 ± 1.56 , 1.64 ± 0.48 , 1.34 ± 0.23 , and 11.5 ± 3.2 μ M, also inhibited aurora-A and B with IC_{50}

values of 90 and 152 nM, respectively. The majority of the compounds with different substitutions at the pyrazine's C-3 position exhibited a range of antiproliferative effects, but compound 8a, which contained 3-chloroaniline in *p*-hydroxybenzoic chains, displayed the strongest inhibitory and antiproliferative activity compared to other compounds. The C-5 position of the pyrazine was changed to 3,5-dimethyl-isoxazole, which also increases the activity of the compound. Compound 8a appears to form stable hydrogen bonds with aurora-A and B (PDB IDs: 3H0Z and 4C2V), according to molecular docking studies. These findings suggested that 8a could be used to generate strong anticancer medicines (**Figure 10**).

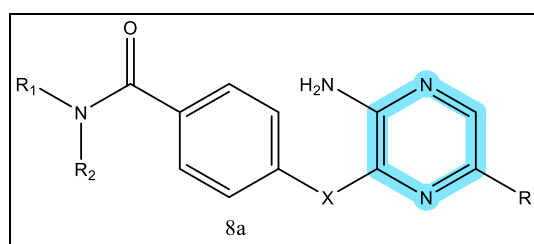


Fig. 10. Structure of compound 8a

Sorafenib kinase inhibitors

Sorafenib is a protein kinase inhibitor that inhibits a variety of kinases, including VEGFR,

PDGFR, and RAF [77, 78]. It targets c-Raf rather than B-RAF. Among the RAF kinases [79], autophagy is induced by sorafenib treatment [80], which may slow tumor growth. It is also a robust soluble epoxide hydrolase inhibitor, thanks to its 1,3-disubstituted urea structure, which likely lessens the severity of its side effects [81].

A variety of sorafenib analogs and derivatives as tumor inhibitors were designed, synthesized, and evaluated [82]. As the lead molecule, sorafenib was employed, with bioisosteres and the alkyl principle being used to achieve changes. Among all, compounds 9a and 9b were shown to be the most powerful and potent with IC₅₀ values of inhibition (3.78±0.21, 16.23±1.23 and 2.34±0.11), and (1.35±0.02, 13.58±1.67 and 2.67±0.12 μmol/l against A549, Hela and H1975 (human lung cancer), respectively. Numerous substituents on the aniline structure were altered during the structural modification of the lead molecule (sorafenib). The following substituents were chosen and tested: the target compounds' physical and chemical characteristics (log P and pKa) improved with the addition of CH₃ at R1, pyrazine and pyridazine at R2, morpholine and pyrimidine at R3, and these substituents also produced high biological activity (**Figure 11**).

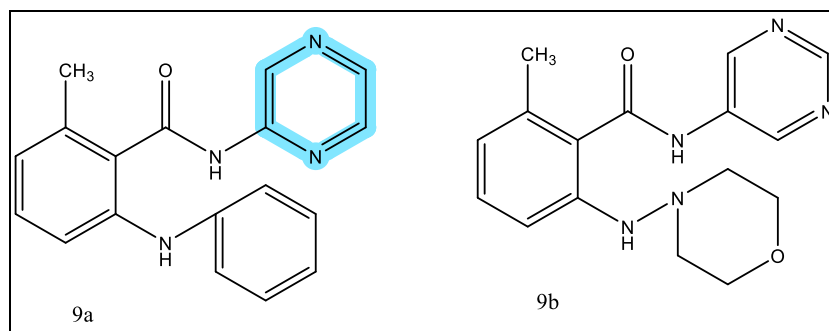


Fig. 11. Structure of compounds 9a and 9b

Miscellaneous

The synthesis of a variety of ligustrazine-chalcone hybrids and tested their anticancer activity [83]. Most of these compounds displayed considerable *in vitro* cytotoxicity against different cell lines. According to the findings, compounds 10a and 10b had the best overall features for antiproliferation effects, with IC₅₀ values of 10a: 1.67±1.25, 1.54±0.30 and 10b: 1.60± 0.21, 1.41± 0.23 μM against MDA-MB231, MCF-7 cell lines, respectively. Furthermore, the addition of the chalcone moiety to the backbone of ligustrazine may

considerably increase its anticancer activity, and the mono substitution of the phenyl moiety at position C-4 in the benzamide segment of the chalcone portion makes it the most potent antiproliferative compound (**Figure 12**).

4-(4-methoxyphenyl)-2-(2-(1-(pyrazin-2-yl) ethylidene) hydrazinyl) thiazole is being used to make iron (II), cobalt (III), and nickel (II) complexes (PyztH). [Fe(Pyzt)₂] Br₂ (Compound 10c), [Co(Pyzt)₂] PF₆ (Compound 10d), and [Ni(Pyzt) (PyztH)] ClO₄ (Compound 10e), having an octahedral environment around the metal center and NNN donor atoms from the two

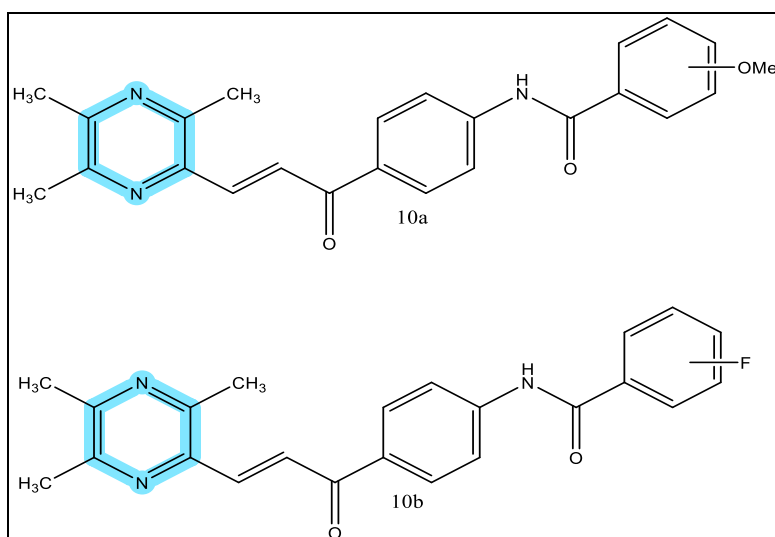


Fig. 12. Structure of compounds 10a and 10b

coordinating ligands [84]. The cytotoxicity of the complexes was tested in U-937 human monocytic cells, with IC_{50} values of 132 (for compound 10c), 45 (compound 10d), and 162

μM (for compound 10e). The docking study also displayed favorable binding of these derivatives with the crystal structure of focal adhesion kinase (FAK) (PDB ID: 6cb0) (**Figure 13**).

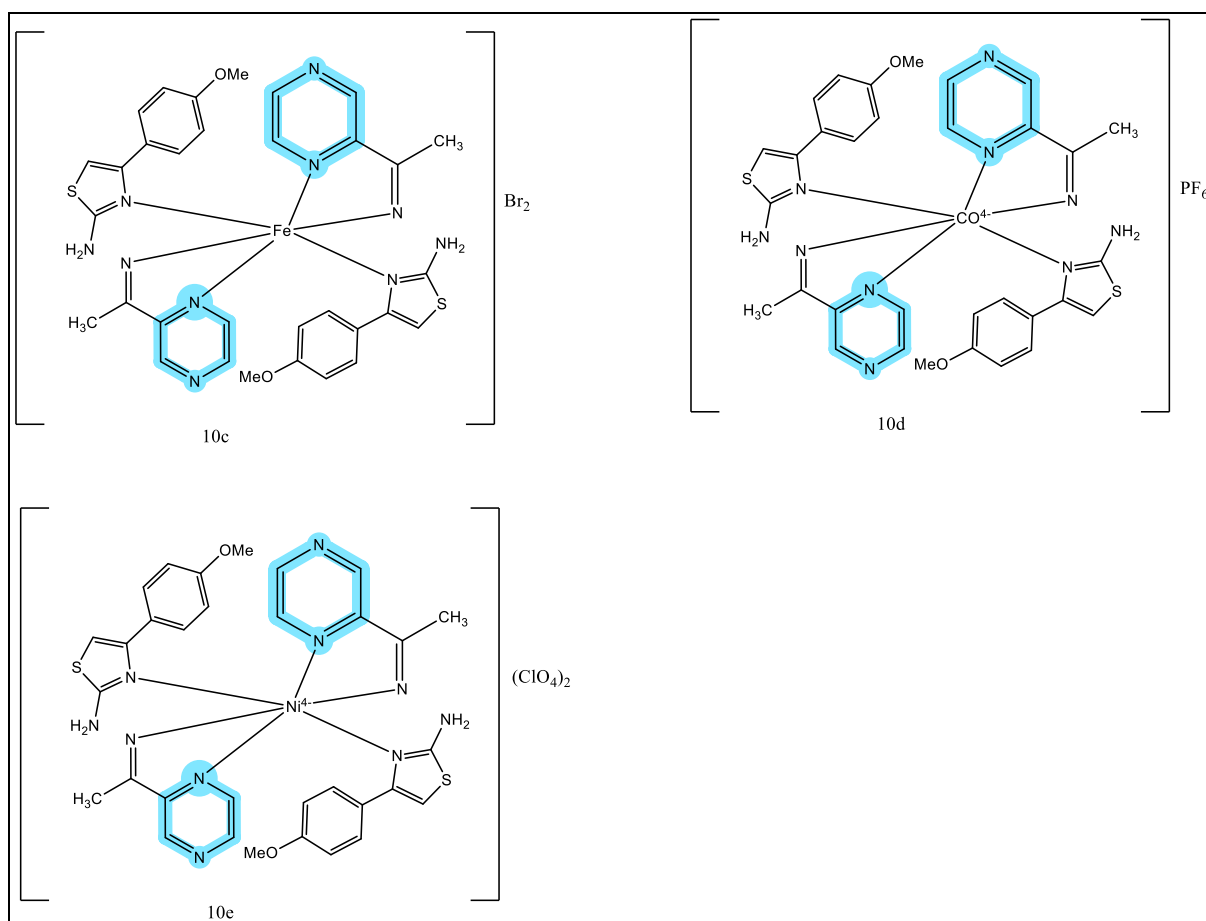


Fig. 13. Structure of compounds 10c-e

The usage of crucial keto-enol intermediates to synthesize two series of A-ring fused steroidal pyrazines from commercially available

progesterone [85]. Additional mechanistic research revealed that the most active molecule, 10f (IC_{50} , 0.93 μM ; SI 28.71), could potentially

cause cell cycle arrest in the G2/M phase and trigger cell death in PC-3 cells in a dose-dependent manner. It seems to fit the active sites of CYP17A1 (PDB ID: 6CIZ) well in a molecular docking analysis. Furthermore, the activities were found to vary significantly depending on where the phenyl ring was replaced; however, beta-phenylethyl substitution displayed the highest level of activity (**Figure 14**).

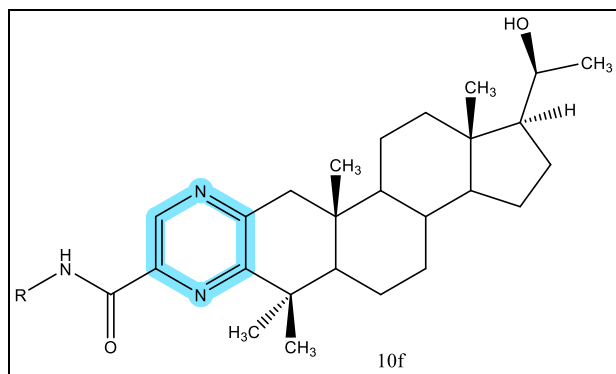


Fig. 14. Structure of compound 10f

Novel hederagenin-pyrazine derivatives, which were tested for *in vitro* cytotoxicity against tumor cell lines [86]. Among all, compound 10g was discovered to be the most powerful molecule, having IC_{50} values of 3.45 ± 0.59 , 8.73 ± 1.49 , 8.71 ± 0.38 , 14.11 ± 0.04 , and 16.69 ± 0.12 μ M against A549, MCF-7, MCF-7, MDCK, and H9c2, respectively. Furthermore, the methyl-substituted pyrazine group of compounds 10g had a significant impact on the anticancer activity; any change or replacement in the pyrazine moiety reduces its activity (**Figure 15**).

CONCLUSION

Following cardiovascular diseases, cancer stands as the second-leading cause of mortality, contributing to one-sixth of global deaths. The rapid developmental strides observed in emerging nations led to a projected 8.9 million

cancer-related deaths in 2016, with an anticipated doubling of these figures by the conclusion of 2040.

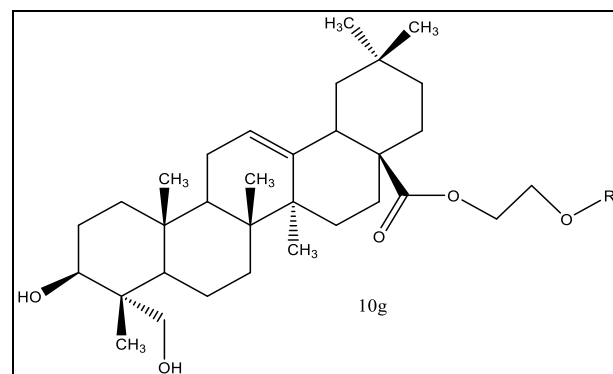


Fig. 15. Structure of compound 10g

Consequently, there is an imperative need for heightened research endeavors directed toward pioneering chemotherapeutic treatments characterized by reduced side effects and heightened efficacy.

This investigation furnishes an abundance of scientific insights aimed at advancing the development of innovative pyrazine analogs or derivatives. These compounds exhibit a diverse array of anticancer characteristics, positioning them within the lucrative and privileged domain of pharmaceutical science. The study delves into various targets associated with these analogs, shedding light on their potential as viable candidates for anticancer therapeutics.

Furthermore, it scrutinizes recently published scientific works across multiple publications, identifying them as prospective alternative treatment options. The comprehensive coverage includes detailed synthesis schemes, tabulated information on current clinical trial drugs, and an overview of recently patented and published research works. This amalgamation of data aims to empower scientists in the development of pharmaceuticals that are not only effective but also possess extraordinary medicinal value.

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