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Development of Pyridazine Derivatives as Potential EGFR inhibitors and Apoptosis Inducers: Design, Synthesis, Anticancer Evaluation, and Molecular Modeling Studies

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Key words: Pyridazine, Pyrazoline; Antiproliferation; epidermal growth factor receptor, Apoptosis, Molecular docking.

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Highlights:

- Novel hybrids of pyridazine-pyrazoline were synthesized aiming to develop new antiproliferative candidates
- All compounds were submitted to the National Cancer Institute (NCI), USA, and many were proved to have significant antiproliferative activity.
- The enzyme inhibitory activity of compounds against epidermal growth factor receptor (EGFR) was evaluated
- A cell cycle analysis and an annexin V-FITC apoptosis assay of the most active compounds was tested.
- all synthesized compounds were subjected to molecular modeling studies using MOE software on the EGFR 3D structure and using Erlotinib as a reference compound.

Abstract

Novel hybrids of pyridazine-pyrazoline were synthesized aiming to develop new antiproliferative candidates. All compounds were submitted to the National Cancer Institute (NCI), USA, and many were proved to have significant antiproliferative activity. In addition, in vitro studies of the epidermal growth factor receptor (EGFR) inhibition showed that compounds IXn, IXg, IXb and IXI exhibited excellent inhibitory effect (IC₅₀=0.65, 0.75, 0.82 and 0.84 µM, respectively) compared to Erlotinib (IC₅₀=0.95 μ M). The mechanistic effectiveness in cell cycle progression, apoptotic induction and gene regulation were assessed for the promising compounds IXg and IXn due to their significant EGFR inhibition. Flow cytometeric analysis indicated that compounds IXg and IXn result in increased cell numbers in phase G2/M, suggesting cell cycle arrest in phase G2/M in UO-31cells. Furthermore, real time PCR assay illustrated that compounds IXg and IXn elevated Bax/Bcl2 ratio which confirmed the mechanistic pathway of them. Moreover, the apoptotic induction of UO-31 renal cancer cells was enhanced effectively through activation of caspase-3 by compounds IXg and IXn. On the other hand, molecular docking study was performed to investigate binding mode of interaction of compounds with EGFR-PK in the active site with the aim of rationalizing its promising inhibitory activity. Finally, based on the aforementioned findings, compounds IXg and IXn could be considered as effective apoptosis modulators and promising leads for future development of new anti-renal cancer agents.

1. Introduction

Despite the rapid advance in drug discovery, the lack of effectiveness, safety and selectivity of anticancer drugs currently available remained as the most important unsolved issue [1]. Targeted therapy, a special type of chemotherapy, is a useful method

for preventing the growth and spread of cancer cells by targeting specific genes or proteins [2]. Advancing our knowledge of molecular and cellular biology enhances our ability to target specific cancer cell functions [3]. The human epidermal growth factor receptor (EGFR) is a tyrosine kinase (RTK) transmembrane receptor consisting of a single polypeptide chain of 1186 amino acids [4, 5]. This receptor belongs to the receptor family ErbB (human epidermal receptor, HER) which comprises four structurally related members EGFR (ErbB1), ErbB2 (HER2), ErbB3 (HER3) and ErbB4 (HER4) [6]. EGFR and members of its family play crucial roles in controlling many cellular processes including cell migration, survival, and proliferation [7]. EGFR may be triggered aberrantly by mutations or over-expression, leading to cell proliferation, angiogenesis, metastasis, and anti-apoptosis. EGFR is over-expressed in a variety of epidermal cancers including head and neck, chest, prostate, breast, cervical, brain and pancreatic cancers [8-10]. Considering that EGFR is a rational target for antitumor strategies, one of the most effective methods of inhibiting EGFR is blocking tyrosine kinase by small-molecule inhibitors at the ATP-binding site in the cytoplasmic domain [6].

In recent years, various nitrogen heterocyclic rings have remarkably complex biological properties and belong to one of the most important classes of medicinal chemistry compounds [11]. Pyrazole motif shows many applications in the field of pharmaceutical and medicinal chemistry. They have anti-inflammatory [12], fungistatic [13], analgesic [14], antibacterial [15], and potent antitumor activities [16-21]. A series of novel compounds containing the pyrazole motifs were reported as potent anticancer agents targeting EGFR tyrosine kinase [22-26]. Furthermore, pyrazoline nucleus represent an important core for triaryl substituted derivatives such as compound 1 (Fig. 1) that had been reported to exhibit powerful cytotoxic activity against different cancer cell lines [27]. In addition, the strategy of hybridization of pyrazoline core with different heterocyclic nuclei such as thiophene (compound 2) [18] and pyrimidine (compound 3) [17] proved effective in production of potent antiproliferative analogs. Therefore, the pyrazoline ring is an advantageous choice for the synthesis of pharmaceutical compounds with diverse biological activities and good safety profiles [28-30]. On the other hand, pyridazine is considered a significant group of heterocyclic compounds which contains two adjacent nitrogen atoms. In recent years, there has been an

increasing interest in the pyridazine ring system due its diverse biological activities and other beneficial applications [31, 32]. Various pyridazine derivatives have been reported to possess increasing attention in the synthesis and evaluation of their biological properties. This indicates that various substituted pyridazine compounds possess different types of pharmacological activities, such as antidiabetic [33], antimicrobial [34], antihypertensive [35], anti-inflammatory [36], and antiproliferative agents [37-40]. Pyridazine **4** (**Fig. 1**) revealed promising cytotoxic activity [41]. A series of novel compounds containing the pyridazine moiety were reported as potent anticancer agents targeting EGFR tyrosine kinase [42]. Finally, from the molecular design point of view, the combination of two pharmacophores into one single molecule represents one of the most important methods that can be used to synthesize new anticancer molecules [43-45].

Fig. 1. Structures of reported antiproliferative compounds with heterocyclic core and triaryl substitution and the structure of the target pyrazoline-pyridazines core hybridization.

This work therefore adopted the hybridization of pyrazoline and pyridazine core with triaryl substitution for the synthesis of a novel pyrazoline/pyridazine hybrids **IXan**, aiming to obtain new promising antiproliferative compounds (**Fig. 1**). The newly synthesized target compounds were sent to the National Cancer Institute (NCI) for assessment of their anticancer activity. Additionally, the most active candidates have

been tested against target EGFR tyrosine kinase inhibition. Also, this study determined annexin V-FITC apoptosis assay. Moreover, the cell cycle activity was detected for the most potent compounds, to get an overview about the possible stage at which the new derivatives could suppress the growth of cancer cells. Furthermore, levels of Bax, Bcl2 and caspase-3 was also determined in this study.

2. Results and discussion

2.1. Chemistry

Compound III was prepared via the reaction of acetophenone I with glyoxilic acid II in the presence of hydrazine hydrate [46]. Furthermore, the reaction between phosphorous oxychloride and compound III afford 3-chloropyridazine IV which has been reacted with hydrazine hydrate to obtain V (Scheme 1) [47].

On the other hand, Chalcones derivatives **VIIIa-n** were prepared by the Claisene Schmidt condensation between acetophenones **VIa** or **VIb** and appropriate aldehyde derivatives **VII** [48-57]. Finally, the target compounds **IXa-n** were prepared by reaction of **V** with the appropriate chalcones **VIII a-n** in absolute ethanol and in the presence of sodium hydroxide (**Scheme 1**). All the final compounds **IXa-n** were characterized by IR, MS, ¹HNMR, ¹³CNMR and elemental analysis.

Scheme 1: Synthesis of compounds IXa-n; Reagents and conditions: (i) hydrazine hydrate, EtOH, reflux, 4h; (ii) phosphorous oxychloride, reflux, 2h; (iii) hydrazine hydrate, EtOH, reflux, 10h.; (iv) ethanolic NaOH, r.t. 6h; (v) ethanolic NaOH, reflux, 15h.

2.2 Biological activity

2.2.1. Screening of anticancer activity and SAR discussion:

Data analysis showed that most synthesized compounds demonstrated different level of growth inhibition against the nine panels of tumor cell lines (**Table 1 and 2**). Furthermore, cell lines of ovarian cancer IGROV1, breast cancer T-47D, renal cancer CAKI-1 and UO-31 were sensitive to almost all evaluated compounds. In this study, we have two series of compounds, first series bearing Cl moiety at position 2 (lead compound **IXa**), the other series bearing methoxy group at position 3 (lead compound **IXh**). Structure–activity correlation was based on the obtained results and in comparison, to the lead compounds, **IXa** and **IXh** by introducing different aryl moieties at 5-position of pyrazole moiety.

The first series, lead compound **IXa** with phenyl moiety at position 5 displayed cell growth inhibition for ovarian cancer IGROV1 (30.19%), renal cancer CAKI-1 (38.28%), renal cancer UO-31 (34.60%), and breast cancer T-47D (41.75%). Substitution with flouro group at position 2 of the phenyl moiety **IXb** increase selectivity against renal cancer CAKI-, renal cancer UO-31, and breast cancer T-47D with growth inhibition (51.05%, 39.54%, 49.22%) respectively.

On the other hand substitution with flouro group at position 3 **IXc** increase selectivity on renal cancer CAKI-1 and UO-31 with cell growth inhibition 40.10%, 46.04%, while substitution with flouro group at position 4 **IXd** shows marked increase growth inhibition against ovarian cancer OVCAR-4 (46.44%), renal cancer CAKI-1 (62.98%), renal cancer UO-31 (51.58%), breast cancer T-47D (55.56%). Compound **IXe** with 3 methyl group at position 3 show only increase in sensitivity toward renal cancer UO-31 (cell growth inhibition: 46.54%). Compound **IXf** with Cl moiety at position 4 decreased in sensitivity toward ovarian cancer IGROV1, breast cancer T-47D, renal cancer CAKI-1, UO-31, cell lines. Furthermore compound **IXg** with flouro group at position 4 and methoxy group at position 3 of the phenyl moiety show marked increase in cell growth inhibition of ovarian cancer IGROV1 (47.17%), renal cancer CAKI-1 (45.86%), renal cancer UO-31 (49.22%).

Table 1: In vitro antiproliferative activity results of compounds IXa - IXg aga	inst sixty
human tumor cell lines; results were provided as a percentage of ce	ll growth
promotion.	

Panel	Subpanel	IXa	IXb	IXc	IXd	IXe	IXf	IXg
	CCRF-CEM	95.17	93.16	98.76	81.77	98.56	98.99	107.00
	HL-60(TB)	98.84	103.32	96.07	92.04	91.22	108.66	95.37
Loukomia	K-562	81.65	88.71	87.85	65.68	81.57	95.98	88.97
Leukenna	MOLT-4	81.48	81.35	88.11	56.77	87.58	94.93	82.30
	RPMI-8226	87.78	86.37	92.61	64.11	85.27	94.73	92.04
	SR	77.36	81.35	87.07	75.76	82.51	85.40	83.78
	A549/ATCC	90.54	96.50	92.13	77.62	88.47	98.96	88.83
	EKVX	90.78	89.85	82.98	77.68	84.38	93.69	79.42
	HOP-62	90.19	89.08	88.15	84.10	94.55	91.65	86.29
Non-small	HOP-92	87.29	78.90	75.41	79.79	79.65	94.24	76.02
cell lung	NCI-H226	87.45	89.73	81.97	74.29	86.45	91.77	77.67
cancer	NCI-H23	95.44	91.38	94.12	77.87	85.96	98.53	86.56
	NCI-H322M	96.73	99.26	97.98	95.68	100.53	103.83	100.12
	NCI-H460	97.59	97.96	98.02	87.15	88.14	99.42	97.57
	NCI-H522	84.59	85.27	84.31	69.25	80.86	96.21	88.20

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	COLO 205	100.76	107.11	105.93	91.34	81.80	104.77	97.59
	HCC-2998	102.88	103 70	106.01	99 42	94 09	98 82	102 10
	HCT-116	81.35	80.60	85.92	57.66	69.17	93 94	80.33
Colon cancer	HCT-15	92.20	90.15	100.53	81.24	91.25	106.80	98.59
	НСТ 10	80.69	91 78	85.12	69.48	58 71	101.87	81.63
	KM12	94 17	99.92	96.77	78 34	92.58	97.67	92.94
	SW-620	97.48	94 98	91.23	85.94	89.23	98.75	89.65
	SF-268	92.58	87.69	91.23	74.16	83.15	92.14	87.11
	SF-295	101.96	99.04	100.12	86.94	101.02	101.28	99 90
	SF-539	93 55	92.61	94 46	83.32	96.72	98.98	95.30
CNS cancer	SNB-19	94.08	90.07	92.93	83.90	88.94	100.18	94.82
	SNB-75	88 73	68.90	86.82	76.41	66.06	65.63	68.24
	U251	89.79	87.90	89.45	77.22	84.36	92.82	90.24
	MALME-3M	91 78	92.61	92 19	86.84	88.58	95.15	94 52
	M14	92.66	91.42	97.55	76.90	87.96	96.19	94.32
	MDA-MB-435	100.91	100 74	101 44	96.33	98.06	101.20	98.94
	SK-MEL-2	101.65	100.74	107.21	93 70	97.30	114 51	102.37
Melanoma	SK-MEL-28	98 77	102.71	107.21	95.62	101.89	106 58	102.57
	SK-MEL-5	85.36	90.05	89.89	78.17	84.12	90.08	79.05
	UACC-257	105.77	100.44	101 79	100.23	98.22	103.78	95.92
	UACC-62	81.50	74 53	78.60	72 37	76.15	89.63	82 50
	IGROV1	69.81	73.86	66.17	58 39	68 46	85.40	52.83
	OVCAR-3	100 31	97.84	98.08	72.38	92.77	97.26	80.63
	OVCAR-4	80.47	81.84	83.29	53 56	79.22	97.20	80.94
Ovarian	OVCAR-5	103 28	96.41	98.22	98.63	100.68	101 42	99 38
cancer	OVCAR-8	93.50	91 45	94 93	80.87	83.56	97.83	89 39
	NCI/ADR-RES	92.60	87.92	94.04	81.18	86.84	101.30	89.19
	SK-OV-3	90.39	96.86	87.85	78.50	83.71	97.73	83.00
	786-0	92.86	97.68	100.17	89.38	89.35	105.45	94.91
	A498	78.45	82.57	90.80	79.34	84.79	103.84	88.26
	ACHN	99.31	94.76	95.97	83.88	95.03	101.48	90.37
D 1	CAKI-1	61.72	48.95	59.90	37.02	60.37	64.66	54.14
Renal cancer	RXF 393	98.54	88.25	98.64	88.06	93.25	109.71	96.45
	SN12C	92.49	91.46	91.61	79.67	92.87	94.68	97.66
	TK-10	103.62	111.44	131.33	130.89	108.60	146.06	151.25
	UO-31	65.40	60.46	53.96	48.42	53.46	67.36	50.78
Prostate	PC-3	76.29	76.87	76.75	56.88	71.43	91.59	71.05
cancer	DU-145	105.04	102.57	106.17	91.83	99.00	109.86	96.85
	MCF7	80.57	82.85	84.78	66.25	80.42	95.27	82.33
	MDA-MB-							
	231/ATCC	89.09	77.05	85.11	75.21	91.55	92.22	91.95
Breast cancer	HS 578T	93.32	93.84	89.59	80.00	94.30	100.42	91.86
	BT-549	95.54	98.34	95.11	87.39	94.01	97.55	95.15
	T-47D	58.25	50.78	73.51	44.44	61.76	69.53	60.67
	MDA-MB-468	96.89	102.35	107.89	79.02	97.37	113.45	99.70

On the other hand, the second series, lead compound **IXh** displayed cell growth inhibition for ovarian cancer IGROV1, renal cancer CAKI-1, renal cancer UO-31 and breast cancer T-47D (47.11%, 33.37%, 41.31%, 20.31%) respectively. Substitution with flouro group at position 2 of the phenyl moiety **IXi** increase selectivity against

renal cancer CAKI-1 (cell growth inhibition: 49.73%), while substitution with flouro group at position 3 **IXj** increase cell growth inhibition of renal cancer UO-31 (47.27%). Furthermore, compounds **IXk** (with flouro moiety at position 4) **IXI** (with methyl moiety at position 3), and **IXm** (with Cl moiety at position 4) show decreased in sensitivity toward ovarian cancer IGROV1, breast cancer T-47D, renal cancer CAKI-1and UO-31, cell lines. Compound **IXn** with flouro group at position 4 and methoxy group at position 3 of the phenyl moiety show increase in cell growth inhibition for renal cancer UO-31 (35.44%), breast cancer T-47D (59.08%).

Table 2: In vitro antiproliferative activity results of compounds IXh - IXn against sixty human tumor cell lines; results were provided as a percentage of cell growth promotion.

Panel	Subpanel	IXh	IXi	IXj	IXk	IXI	IXm	IXn
	CCRF-CEM	99.59	95.14	96.27	99.70	96.26	90.56	94.47
Leukemia	HL-60(TB)	101.11	104.71	86.76	98.66	103.81	87.37	97.89
	K-562	92.60	88.84	83.25	96.80	87.33	84.32	84.03
	MOLT-4	90.64	76.66	80.54	96.78	86.48	82.54	82.21
	RPMI-8226	97.99	87.99	81.10	104.50	89.85	78.08	80.57
	SR	95.27	82.32	78.32	86.70	82.86	79.14	84.67
	A549/ATCC	93.99	88.73	85.57	96.99	94.54	91.02	95.14
	EKVX	NT	76.41	83.81	91.46	82.79	83.26	86.73
	HOP-62	79.40	81.98	95.20	91.27	89.82	90.71	99.67
NT II II	HOP-92	77.68	70.54	75.85	87.34	85.08	91.47	93.01
Non-small cell	NCI-H226	71.14	74.42	82.55	89.69	85.03	85.24	92.93
lung cancer	NCI-H23	90.40	86.20	89.19	92.08	85.75	82.20	89.90
	NCI-H322M	85.48	88.64	92.34	92.98	91.18	96.54	101.50
	NCI-H460	97.28	94.38	97.79	99.28	97.71	88.38	96.79
	NCI-H522	93.49	86.51	80.65	94.34	89.29	83.83	91.48
	COLO 205	106.93	107.55	94.30	107.01	99.73	88.30	96.53
	HCC-2998	98.98	105.04	98.51	100.56	103.84	97.28	105.27
	HCT-116	80.09	80.57	82.45	100.80	81.78	68.22	68.24
Colon cancer	HCT-15	NT	94.95	85.39	100.15	88.59	85.74	91.02
	HT29	100.76	93.14	83.66	102.06	91.00	67.28	82.86
	KM12	96.87	87.20	92.78	98.68	97.31	89.32	93.00
	SW-620	93.34	89.46	90.80	100.04	92.62	82.00	92.26
	SF-268	86.68	88.64	85.33	93.75	90.92	87.19	91.28
	SF-295	NT	95.42	101.51	101.06	95.61	100.22	100.24
CNS concor	SF-539	93.49	94.57	91.44	94.45	94.79	95.74	96.31
CNS cancer	SNB-19	87.63	87.70	82.17	91.94	87.85	87.68	92.84
	SNB-75	91.83	78.85	82.99	83.87	98.18	78.34	103.49
	U251	91.13	90.61	86.21	97.04	92.25	84.53	90.87
	MALME-3M	84.38	93.28	88.39	97.34	95.19	93.94	100.96
	M14	87.93	91.33	88.74	102.05	88.90	85.99	94.59
Melanoma	MDA-MB- 435	101.22	100.95	99.98	102.04	100.83	97.92	100.87
	SK-MEL-2	102.59	98.46	83.75	105.21	97.46	94.64	98.06

	SK-MEL-28	99.68	105.33	98.38	103.04	101.89	98.85	106.00
	SK-MEL-5	88.62	91.86	89.43	98.36	98.22	94.52	84.86
	UACC-257	96.56	98.35	96.58	108.24	99.92	98.01	104.54
	UACC-62	75.70	73.07	77.23	94.59	75.52	79.65	86.96
	IGROV1	52.89	61.59	57.14	89.38	72.10	72.62	85.75
	OVCAR-3	91.33	91.72	89.90	106.96	99.90	96.92	100.73
	OVCAR-4	93.48	82.61	84.39	99.68	85.46	89.54	76.78
Ovarian cancer	OVCAR-5	95.02	98.16	92.50	99.58	90.35	96.34	100.07
Ovarian cancer	OVCAR-8	94.23	93.71	87.66	99.54	93.71	89.08	93.38
	NCI/ADR- RES	92.91	94.32	86.23	101.68	95.01	93.79	95.24
	SK-OV-3	84.91	89.75	85.32	96.28	92.99	79.34	100.10
	786-0	102.02	106.26	86.19	99.25	100.25	89.56	98.43
	A498	57.41	78.32	66.28	79.22	76.01	75.27	76.08
	ACHN	87.47	87.51	90.98	98.65	97.95	95.81	101.58
Donal concor	CAKI-1	66.63	50.27	68.52	79.54	65.27	71.06	73.33
Kenai cancer	RXF 393	91.73	102.26	85.00	96.96	92.08	98.11	110.69
	SN12C	91.35	90.47	89.07	93.20	88.15	98.59	101.32
	TK-10	145.99	170.13	131.49	125.03	143.16	137.87	109.39
	UO-31	58.69	62.38	52.73	85.86	75.67	67.82	64.56
Prostata gangar	PC-3	84.53	79.65	73.51	100.02	85.93	71.20	78.78
1 I Ostate cancer	DU-145	101.12	102.87	100.08	102.87	98.84	99.93	102.32
	MCF7	NT	83.96	79.82	93.95	82.71	76.22	85.62
	MDA-MB- 231/ATCC	74.15	78.02	86.48	86.96	77.70	85.54	97.06
Durant comon	HS 578T	94.46	81.30	88.48	95.79	95.78	85.05	94.14
breast cancer	BT-549	88.41	91.37	93.30	96.65	94.15	91.63	97.45
	T-47D	79.69	74.00	77.44	99.94	85.94	72.82	40.92
	MDA-MB- 468	89.67	97.15	92.35	100.47	99.54	99.50	95.35

NT: Not tested.

2.2.2. IC₅₀ determination of the synthesized compounds against EGFR

All new synthesized compounds **IXa-n** were further investigated for their inhibitory effect on human EGFR using Erlotinib as a positive control to determine their IC₅₀ values as shown in (**Table 3**). The tested compounds showed significant inhibitory effect (IC₅₀=5.5–0.65 μ M). It was found that compounds **IXn**, **IXg**, **IXb** and **IXI** inhibition effect more than Erlotinib towards EGFR (IC₅₀=0.65, 0.75, 0.82 and 0.84 μ M, respectively) compared to Erlotinib (IC₅₀=0.95 μ M). Compound **IXc** had the same potency as Erlotinib against EGFR.

 Table 3:
 Inhibition activities of compounds IXh - IXn against EGFR.

Comp.	EGFR IC ₅₀ (µM)
IXa	2.1 ± 0.12
IXb	0.82 ± 0.06
IXc	0.95 ± 0.07

IXd	3.7 ± 0.26
IXe	3.9 ± 0.2
IXf	4.8 ± 0.25
IXg	0.75 ± 0.05
IXh	1.7 ± 0.14
IXi	5.5 ± 0.41
IXj	4.1 ± 0.3
IXk	5.3 ± 0.43
IXI	0.84 ± 0.06
IXm	3.1 ± 0.23
IXn	0.65 ± 0.03
Erlotinib	0.95 ± 0.08

2.2.3. Cell cycle analysis

Results in **Tables 1** and **2** revealed that compounds **IXg** and **IXn** exhibited potent cytotoxic activity towards renal cancer cell line UO-31. In this manner, the effect of both compounds on the cell cycle progression of UO-31 cancer cells were performed to carry out preliminary mechanistic studies of their anticancer activity. The cancer cells were incubated for 24 h with compounds **IXg** and **IXn** at 10 μ M. Results analysis indicated that both compounds resulted in increased of cell numbers in phase G2/M (50.11 % vs. 45.26 % and 12.39 %), suggesting that cell cycle was arrested in phase G2/M (**Table 4, Fig. 2**).

 Table 4: Effect of compounds IXg and IXn on the cell cycle progression of UO-31 cells.

Code	%G0/G1	%S	%G2/M	%Pre-G1	Comment
IXg/ UO-31	24.45	25.44	50.11	20.17	Cell growth arrest@G2/M
IXn/ UO-31	28.14	26.6	45.26	15.36	Cell growth arrest@G2/M
Cont. (UO-31)	51.28	36.33	12.39	1.85	

IXn

2.2.4. Annexin V-FITC apoptosis assay

Cell cycle assay of UO-31 cancer cells treated with compounds **IXg** and **IXn** showed pre-G1 peak appearance which emphasized apoptosis induction. The cells were stained with Annexin V / PI, incubated for 24 h and analyzed to confirm the efficacy of these compounds to cause apoptosis. It has been found that the early and late apoptosis induced by the compounds being studied definitely suggested their ability to cause meaningful apoptosis rates with 1.5 and 2.7 % necrosis respectively (**Table 5, Fig. 3**). In addition, the percentages of late apoptosis (10.29% and 7.25%) induced by **IXg** and **IXn** were higher than those of early apoptosis (8.37% and 5.41%) which makes recovery of apoptotic cells more difficult to be healthy.

Table 5. Induced apoptosis and necrosis on UO-31 cells by IXg and IXn.

		Neerosia		
	Total	Early	Late	INECTOSIS
IXg/UO-31	20.17	8.37	10.29	1.51
IXn/UO-31	15.36	5.41	7.25	2.7
Cont. (UO-31)	1.85	1.13	0.29	0.43

Fig. 3. Effect of compounds IXg and IXn on the percentage of V-FITC-positive annexin staining in UO-31 cells.

2.2.5. Effects on the levels of Bax, Bcl-2 and active caspase-3

Apoptosis is regulated by various factors including stimulation of the apoptotic pathway through caspase-3, inducers (Bax protein) and suppressors (Bcl-2 protein) [58]. The key members of apoptotic pathways are the cysteine protease caspase family [59]. Caspase-3 is an apoptosis executioner and that its activation contributes to apoptosis induction [58]. Many studies on human cancers have revealed a relationship between increased Bcl-2 levels, decreased Bax levels and uncontrolled cell growth [59, 60]. Therefore, in this study, the levels of Bcl-2, Bax and caspase-3 were determined after treatment with **IXg** and **IXn** in UO-31 cells in order to examine the effect of both compounds on these apoptotic biomarkers. The findings presented in (**Table 6**) showed that **IXg** and **IXn** raised the Bax level by 7.3 and 6.6 folds, respectively, and decreased the Bcl-2 level by 0.31 and 0.44 folds, respectively, compared with the control. Also results showed that the active caspase-3 was significantly overexpressed by 7.1 and 5.9 folds relative to the control indicating the apoptosis induction potential of these compounds.

Comp.	Bax conc. pg	FLD	Bcl-2 Conc. pg	FLD	Caspase-3 conc. pg	FLD
13	392.17	7.33	1.84	0.317	419.25	7.19
14	356.25	6.6	2.56	0.44	347.11	5.95
Cont.	53.49	1	5.79	1	58.31	1

 Table 6. Bax, Bcl2 and Caspase-3 analysis results after treatment with compound IXg and IXn.

2.3. Molecular docking

Molecular docking studies are an important method for studying molecular interactions between the synthesized ligands and main amino acids at the respective protein binding site [61]. Ligand – protein interaction activity at the binding active site was measured on the basis of the docking score values determined using MOE 2015.10. In the present research, all synthesized compounds were subjected to molecular modeling studies using MOE software on the EGFR 3D structure and using Erlotinib as a reference compound Table 7, in order to predict the behavior of the synthesized compounds as an anticancer agents. The results of the docking studies showed that the binding free energy of the compounds being examined exhibited favorable docking complexes with the active target protein site. The compounds being studied displayed interactions with the active EGFR sites with the same amino acid as did the reference drug, Erlotinib. Although, all the tested compounds showed significant docking score free energy compared to Erlotinib, compounds **IXc**, **IXg**, **IXj**, **IXe**, and **IXn** showed the best docking score free energy (**Table 7**).

Comp. No.	Score	rmsd_ refine	E_conf	E_place	E_score1	E_refine	E_score2
IXa	-7.5032	1.1158	52.5396	-67.4750	-13.0213	-43.1217	-7.5032
IXb	-7.4828	1.1512	52.4754	-82.2369	-12.2609	-35.5909	-7.4828
IXc	-8.0715	1.5914	46.3532	-53.7557	-11.9093	-42.3909	-8.0715
IXd	-7.5669	1.2320	52.2871	-76.9072	-13.2147	-41.1207	-7.5669
IXe	-7.6098	1.1562	50.0204	-99.3674	-11.9157	-40.4771	-7.6098
IXf	-7.5063	2.3522	49.8316	-71.1179	-12.5182	-43.0492	-7.5063
IXg	-7.7437	1.2496	70.0082	-93.1728	-12.3985	-37.4756	-7.7437
IXh	-7.4173	2.0378	61.9342	-66.4482	-12.0435	-40.5859	-7.4173

 Table 7. Docking energy scores (kcal/mol) derived from the MOE for Compounds IXan and Erlotinib.

IXi	-7.4513	1.1826	55.3965	-67.8661	-11.5407	-38.7661	-7.4513
IXj	-7.6264	1.1773	62.3024	-58.6817	-12.0223	-41.9693	-7.6264
IXk	-7.4602	1.9206	64.0970	-66.7235	-12.1795	-40.9835	-7.4602
IXI	-7.4967	1.9904	62.7921	-66.7189	-11.4621	-39.5500	-7.4967
IXm	-7.4797	1.3109	55.0005	-86.0764	-13.1680	-38.6300	-7.47977
IXn	-7.5941	1.8792	78.2140	-68.6826	-12.3753	-40.3379	-7.5941
Erlotinib	-7.3430	2.9191	-34.3322	-71.5678	-12.4086	-42.4795	-7.3430

Score; lower scores are more favorable, **rmsd_refine**; the root mean square deviation of the pose from the docking pose compared to the co-crystal ligand position, **E_conf**; free binding energy of the conformer, **E_place**; free binding energy from the placement stage, **E_score 1**; free binding energy from the first rescoring stage, **E_refine**; free binding energy from the refinement stage, **E_score 2**; free binding energy from the second rescoring stage.

Compound IXg and IXn showed an excellent docking score (-7.7437 kcal/mol and -7.5941 kcal/mol) compared to Erlotinib docking score (-7.3430 kcal/mol), and both formed a direct interaction with most of the amino-acids that Erlotinib interacted with. Where, Erlotinib had a direct interaction with amino-acids VAL702, LYS721, CYS751 and THR766 in the active site (Fig. 4A), compound IXg and IXn showed interactions with amino-acids LEU694, LYS721 and THR766 in the active site (Fig4 4B and 4C). Also, compounds IXg and IXn showed high degrees of superimposition Erlotinib into EGFR active sites (Fig. 4B and 4C). These results were in a perfect match with the EGFR enzyme inhibition assay where compounds IXg and IXn had IC_{50s} of (0.75 μ M/mL and 0.65 μ M/mL, respectively) compared to that of Erlotinib (0.95 μ M/mL) (Table 3).

A) Interaction of Erlotinib with amino-acids VAL702, LYS721, CYS751 and THR766.

B) Interaction of **IXg** with amino-acids LEU694, LYS721 and THR766 and superimposition of **IXg** (shown as grey sticks) with **Erlotinib** (shown as cyan sticks).

C) Interaction of **IXn** with amino-acids LEU694, LYS721 and THR766 and superimposition of **IXn** (shown as grey sticks) with **Erlotinib** (shown as cyan sticks).

Fig. 4. Docking of compounds IXg, IXn and Erlotinib into EGFR active sites.

3. Experimental

3.1. Chemistry

IR spectra had been measured on a Perkin Elmer-9712 spectrophotometer. Elemental microanalysis was performed at the Micro-analytical Laboratory at Cairo University, Egypt. ¹H NMR spectrometer and ¹³C NMR spectra were measured on Bruker 400 MHz NMR. For the evaluation of mass spectra, a Shimadzu Qp-2010 Plus was used. 6-phenylpyridazines V [46, 47] and Chalcones VIII [48-57] have been prepared according to the procedures indicated.

3.1.1. General procedure for preparation of IXa-n

In absolute ethanol (15 mL), a mixture of the corresponding chalcones **VIIIa-n** (1 mmol), hydrazine derivatives **V** (1 mmol) and sodium hydroxide (2.5 mmol) were heated for 15 h under reflux. Filtered and washed with water and crystallized with ethanol.

3.1.1.1. 3-(3-(2-chlorophenyl)-4,5-dihydro-5-phenylpyrazol-1-yl)-6-phenylpyridazine (IXa)

White powder (yield 64%); m.p. 107-109 °C; IR (v_{max} /cm⁻¹) 3031 (C-H aromatic), 1649 (C=N) and at 1588 (C=C); ¹HNMR (400 MHz, DMSO-d₆) δ *ppm*: 8.08 (d, J = 9.3 Hz, 1H, Ar-H), 8.00 (d, J = 7.6 Hz, 2H, Ar-H), 7.81-7.85 (m, 1H, Ar-H), 7.78 (d, J = 9.5 Hz, 1H, Ar-H), 7.56-7.60 (m, 1H, Ar-H), 7.41-7.50 (m, 6H, Ar-H), 7.30-7.36 (m, 4H, Ar-H), 5.94 (dd, J = 5.1, 12.0 Hz, 1H, pyrazoline-CH), 4.16 (dd, J = 12.0, 17.9 Hz, 1H, one proton of pyrazoline-CH₂), 3.31 (d, J = 5.4 Hz, 1H, one proton of pyrazoline-CH₂); ¹³CNMR (100 MHz, DMSO-d₆) δ *ppm*: 155.9, 152.4, 151.1, 142.9, 136.8, 131.9, 131.4, 131.2, 131.1, 130.7, 129.4, 129.3, 129.2, 127.9, 127.7, 126.7, 126.3, 126.2, 115.4, 62.1, 45.7; MS (m/z (R.I. %)); [M]⁺ 410 (12.7), [M+2]⁺ 412 (4.7), 272 (100), 170 (17.4); Anal. Calcd. for C₂₅H₁₉ClN₄ (410.13); % C, 73.08; H, 4.66; N, 13.64, Found: % C, 73.21; H, 4.75; N, 13.71.

3.1.1.2. 3-(3-(2-chlorophenyl)-5-(2-fluorophenyl)-4,5-dihydropyrazol-1-yl)-6phenyl-pyridazine (IXb)

White powder (yield 60%); m.p. Over 280 °C; IR (v_{max} /cm⁻¹) 3054 (C-H aromatic), 1649 (C=N) and at 1589 (C=C); ¹HNMR (400 MHz, DMSO-d₆) δ *ppm*: 8.10 (d, J = 9.5 Hz, 1H, Ar-H), 8.00 (d, J = 7.6 Hz, 2H, Ar-H), 7.82-7.86 (m, 1H, Ar-H), 7.79 (d, J = 9.3 Hz, 1H, Ar-H), 7.55-7.61 (m, 1H, Ar-H), 7.42-7.50 (m, 5H, Ar-H), 7.29-7.34 (m, 1H, Ar-H), 7.20-7.28 (m, 2H, Ar-H), 7.13 (t, J = 7.3 Hz, 1H, Ar-H), 6.09 (dd, J = 5.6, 12.2 Hz, 1H, pyrazoline-CH), 4.19 (dd, J = 12.4, 17.7 Hz, 1H, one proton of pyrazoline-CH₂), 3.30 (m, 1H, one proton of pyrazoline-CH₂); MS (m/z (R.I. %)); [M]⁺ 428 (25.5), [M+2]⁺ 430 (9.7), 290 (79.5), 272 (100), 170 (31.2); Anal. Calcd. For C₂₅H₁₈ClFN₄ (428.12); % C, 70.01; H, 4.23; N, 13.06, Found: % C, 70.09; H, 4.35; N, 13.15.

3.1.1.3. 3-(3-(2-chlorophenyl)-5-(3-fluorophenyl)-4,5-dihydropyrazol-1-yl)-6phenylpyridazine (IXc)

Yellowish powder (yield 59%); m.p. 209-211 °C; IR (v_{max}/cm^{-1}) 3064 (C-H aromatic), 1647 (C=N) and at 1590 (C=C); ¹H NMR (400 MHz, DMSO-d₆) δ *ppm*: 8.10 (d, *J* = 8.8 Hz, 1H, Ar-H), 7.97-8.04 (m, 2H, Ar-H), 7.77-7.87 (m, 2H, Ar-H), 7.57 (br. s, 1H, Ar-H), 7.44-748 (m, 6H), 7.16 (br. s, 2H, Ar-H), 7.06-7.13 (m, 1H, Ar-H), 5.87-6.03 (m, 1H, pyrazoline-CH), 4.05-4.24 (m, 1H, one proton of pyrazoline-CH₂), 3.32

(m, 1H, one proton of pyrazoline-CH₂); MS (m/z (R.I. %)); [M-F]⁺ 409 (31.5), 386 (20.7), 306 (43.1), 55 (100); Anal. Calcd. for C₂₅H₁₈ClFN₄ (428.12); % C, 70.01; H, 4.23; N, 13.06, Found: % C, 70.12; H, 4.19; N, 12.98.

3.1.1.4. 3-(3-(2-chlorophenyl)-5-(4-fluorophenyl)-4,5-dihydropyrazol-1-yl)-6phenyl-pyridazine (IXd)

Yellowish powder (yield 63%); m.p. 183-185 °C; IR (v_{max}/cm^{-1}) 3038 (C-H aromatic), 1649 (C=N) and at 1588 (C=C); ¹HNMR (400 MHz, DMSO-d₆) δ *ppm*: 8.07 (d, *J* = 9.5 Hz, 1H, Ar-H), 8.00 (d, *J* = 7.3 Hz, 2H, Ar-H), 7.80-7.86 (m, 1H, Ar-H), 7.77 (d, *J* = 9.5 Hz, 1H, Ar-H), 7.54-7.60 (m, 1H, Ar-H), 7.41-7.50 (m, 5H, Ar-H), 7.36 (dd, *J* = 5.5, 8.0 Hz, 2H, Ar-H), 7.16 (t, *J* = 8.8 Hz, 2H, Ar-H), 5.94 (dd, *J* = 5.4, 12.0 Hz, 1H, pyrazoline-CH), 4.14 (dd, *J* = 12.2, 17.9 Hz, 1H, one proton of pyrazoline-CH₂), 3.31-3.37 (m, 1H, one proton of pyrazoline-CH₂); ¹³CNMR (100 MHz, DMSO-d₆) δ *ppm*: 155.9, 152.5, 151.1, 136.7, 131.9, 131.4, 131.3, 131.1, 130.6, 129.4, 129.3, 128.4, 128.3, 128.0, 126.3, 126.2, 116.1, 115.8, 115.5, 61.5, 45.6; MS (m/z (R.I. %))); [M]⁺ 428 (12.9), [M+2]⁺ 430 (4.7), 290 (100), 170 (15.5), 115 (35.3); Anal. Calcd. for C₂₅H₁₈ClFN₄ (428.12); % C, 70.01; H, 4.23; N, 13.06, Found: % C, 70.09; H, 4.27; N, 13.01.

3.1.1.5. 3-(3-(2-chlorophenyl)-4,5-dihydro-5-m-tolylpyrazol-1-yl)-6-phenylpyridazine (IXe)

Buff powder (yield 55%); m.p. Over 280 °C; IR (v_{max}/cm^{-1}) 3050 (C-H aromatic), 1648 (C=N) and at 1590 (C=C); ¹HNMR (400 MHz, DMSO-d₆) δ *ppm*: 8.08 (d, *J* = 9.3 Hz, 1H, Ar-H), 8.00 (d, *J* = 6.9 Hz, 2H, Ar-H), 7.83 (br. s, 1H, Ar-H), 7.78 (d, *J* = 9.3 Hz, 1H, Ar-H), 7.54-7.61 (m, 1H, Ar-H), 7.41-7.51 (m, 5H, Ar-H), 7.18-7.25 (m, 1H, Ar-H), 7.15 (br. s, 1H, Ar-H), 7.07 (t, *J* = 8.2 Hz, 2H, Ar-H), 5.83-5.95 (m, 1H, pyrazoline-CH), 4.15 (dd, *J* = 12.4, 17.2 Hz, 1H, one proton of pyrazoline-CH₂), 3.30 (br. s, 1H, one proton of pyrazoline-CH₂), 2.28 (s, 3H, CH₃); ¹³CNMR (100 MHz, DMSO-d₆) δ *ppm*: 155.9, 152.4, 151.1, 143.0, 138.4, 136.7, 131.9, 131.4, 131.2, 131.1, 130.7, 129.3, 129.2, 128.4, 128.0, 126.7, 126.3, 126.2, 123.1, 115.4, 62.1, 45.8, 21.5; MS (m/z (R.I. %)); [M]⁺ 424 (9.7), [M+2]⁺ 426 (3.6), 286 (100), 170 (14.6), 115 (37.3)); Anal. Calcd. for C₂₆H₂₁ClN₄ (424.15); % C, 73.49; H, 4.98; N, 13.19, Found: % C, 73.55; H, 4.87; N, 13.25.

3.1.1.6. 3-(3-(2-chlorophenyl)-5-(4-chlorophenyl)-4,5-dihydropyrazol-1-yl)-6phenylpyridazine (IXf)

Yellowish powder (yield 60%); m.p. Over 280 °C; IR (v_{max}/cm^{-1}) 3059 (C-H aromatic), 1677 (C=N) and at 1587 (C=C); ¹HNMR (400 MHz, DMSO-d₆) δ *ppm*: 8.10 (d, J = 9.5 Hz, 1H, Ar-H), 8.00 (d, J = 7.6 Hz, 2H, Ar-H), 7.81-7.86 (m, 1H, Ar-H), 7.78 (d, J = 9.3 Hz, 1H, Ar-H), 7.35-7.42 (m, 6H, Ar-H), 7.11 (d, J = 8.1 Hz, 2H, Ar-H), 6.96 (d, J = 8.8 Hz, 2H, Ar-H), 5.94 (dd, J = 5.5, 11.9 Hz, 1H, pyrazoline-CH), 4.16 (dd, J = 12.2, 17.6 Hz, 1H, one proton of pyrazoline-CH₂); ¹³CNMR (100 MHz, DMSO-d₆) δ *ppm*: 155.9, 152.5, 151.2, 141.9, 138.6, 136.7, 136.3, 132.8, 131.9, 131.4, 131.2, 130.0, 129.3, 129.2, 128.3, 127.9, 126.4, 126.2, 115.4, 61.6, 45.5; MS (m/z (R.I. %)); [M]⁺ 444 (9.8), [M+2]⁺ 446 (6.7), 306 (100), 308 (37), 115 (50.2); Anal. Calcd. for C₂₅H₁₈Cl₂N₄ (444.09); % C, 67.42; H, 4.07; N, 12.58, Found: % C, 67.40; H, 3.93; N, 12.65.

3.1.1.7. 3-(3-(2-chlorophenyl)-5-(4-fluoro-3-methoxyphenyl)-4,5-dihydropyrazol-1yl)-6-phenylpyridazine (IXg)

Buff powder (yield 64%); m.p. 166-168 °C; IR (v_{max}/cm^{-1}) 3059 (C-H aromatic), 1648 (C=N) and at 1589 (C=C); ¹HNMR (400 MHz, DMSO-d₆) δ *ppm*: 8.09 (d, *J* = 9.5 Hz, 1H, Ar-H), 7.99 (d, *J* = 7.6 Hz, 2H, Ar-H), 7.81-7.86 (m, 1H, Ar-H), 7.73-7.76 (m, 2H, Ar-H), 7.33-7.37 (m, 4H, Ar-H), 7.06-7-09 (m, 2H, Ar-H), 6.94 (d, *J* = 8.8 Hz, 2H, Ar-H), 5.92 (dd, *J* = 5.6, 12.2 Hz, 1H, pyrazoline-CH), 4.14 (dd, *J* = 12.2, 17.6 Hz, 1H, one proton of pyrazoline-CH₂), 3.84 (s, 3H, OCH₃), 3.33 (d, *J* = 5.6 Hz, 1H, one proton of pyrazoline-CH₂); MS (m/z (R.I. %)): [M]⁺ 458 (13.5), [M+2]⁺ 460 (5.4), 320 (100), 197 (15.2), 115 (27); Anal. Calcd. for C₂₆H₂₀ClFN₄O (458.13); % C, 68.05; H, 4.39; N, 12.21, Found: % C, 67.96; H, 4.42; N, 12.16.

3.1.1.8. 3-(4,5-dihydro-3-(3-methoxyphenyl)-5-phenylpyrazol-1-yl)-6-phenylpyridazine (IXh)

Buff powder (yield 60%); m.p. 162-164 °C; IR (v_{max}/cm^{-1}) 2960 (C-H aromatic), 1668 (C=N) and at 1597 (C=C); ¹HNMR (400 MHz, DMSO-d₆) δ *ppm*: 8.06 (d, *J* = 8.8 Hz, 1H, Ar-H), 8.01 (d, *J* = 7.6 Hz, 2H, Ar-H), 7.80-7.84 (m, 2H, Ar-H), 7.62 (br. s, 1H, Ar-H), 7.29-7.44 (m, 6H, Ar-H), 7.16 (br. s, 2H, Ar-H), 7.01-7.8 (m, 2H, Ar-H), 5.92 (dd, J = 5.6, 12.2 Hz, 1H, pyrazoline-CH), 3.99 (dd, J = 12.2, 17.9 Hz, 1H, one proton of pyrazoline-CH₂), 3.82 (s, 3H, OCH₃), 3.31 (dd, J = 5.6, 17.9 Hz, 1H, one proton of pyrazoline-CH₂); MS (m/z (R.I. %)): [M]⁺ 406 (22.5), 272 (100), 170 (14.3), 115 (26); Anal. Calcd. for C₂₆H₂₂N₄O (406.18); % C, 76.83; H, 5.46; N, 13.78, Found: % C, 76.75; H, 5.42; N, 13.69.

3.1.1.9. 3-(5-(2-fluorophenyl)-4,5-dihydro-3-(3-methoxyphenyl)pyrazol-1-yl)-6phenylpyridazine (IXi)

Yellowish powder (yield 56%); m.p. 123-125 °C; IR (v_{max}/cm^{-1}) 3060 (C-H aromatic), 1647 (C=N) and at 1596 (C=C); ¹HNMR (400 MHz, DMSO-d₆) δ *ppm*: 8.08 (d, J = 9.3 Hz, 1H, Ar-H), 8.00 (d, J = 7.8 Hz, 2H, Ar-H), 7.86 (d, J = 9.3 Hz, 1H, Ar-H), 7.77-7.81 (m, 1H, Ar-H), 7.63 (br. s, 1H, Ar-H), 7.31-7.44 (m, 5H, Ar-H), 7.17-24 (m, 3H, Ar-H), 7.03-7.6 (m, 1H, Ar-H), 6.03 (dd, J = 5.5, 12.1 Hz, 1H, pyrazoline-CH), 4.04 (dd, J = 12.2, 17.9 Hz, 1H, one proton of pyrazoline-CH₂); MS (m/z (R.I. %)): [M]⁺ 424 (35.3), 290 (80.4), 272 (100), 170 (30), 115 (44.2); Anal. Calcd. for C₂₆H₂₁FN₄O (424.17); % C, 73.57; H, 4.99; N, 13.20, Found: % C, 73.49; H, 5.12; N, 13.13.

3.1.1.10. 3-(5-(3-fluorophenyl)-4,5-dihydro-3-(3-methoxyphenyl)pyrazol-1-yl)-6phenylpyridazine (IXj)

Yellowish powder (yield 55%); m.p. 177-179 °C; IR (v_{max} /cm⁻¹) 3082 (C-H aromatic), 1649 (C=N) and at 1593 (C=C); ¹HNMR (400 MHz, DMSO-d₆) δ *ppm*: 8.08 (d, J = 9.5 Hz, 1H, Ar-H), 8.01 (d, J = 7.6 Hz, 2H, Ar-H), 7.86 (d, J = 9.3 Hz, 1H, Ar-H), 7.36-7.51 (m, 7H, Ar-H), 7.01-7.16 (m, 4H, Ar-H), 5.94 (dd, J = 5.5, 12.1 Hz, 1H, pyrazoline-CH), 4.01 (dd, J = 12.2, 17.9 Hz, 1H, one proton of pyrazoline-CH₂), 3.83 (s, 3H, OCH₃), 3.30 (dd, J = 5.6, 17.9 Hz, 1H, one proton of pyrazoline-CH₂); ¹³CNMR (100 MHz, DMSO-d₆) δ *ppm*: 161.6, 159.9, 155.9, 152.6, 152.3, 146.1, 136.8, 133.2, 131.2, 130.4, 129.3, 126.2, 122.2, 119.4, 116.3, 115.4, 114.5, 114.3, 113.3, 111.7, 61.8, 55.7, 43.1; MS (m/z (R.I. %)): [M]⁺ 424 (21.9), 290 (100), 170 (16.2), 115 (33.7); Anal. Calcd. for C₂₆H₂₁FN₄O (424.17); % C, 73.57; H, 4.99; N, 13.20, Found: % C, 73.62; H, 4.94; N, 13.29.

3.1.1.11. 3-(5-(4-fluorophenyl)-4,5-dihydro-3-(3-methoxyphenyl)pyrazol-1-yl)-6phenylpyridazine (IXk).

White powder (yield 63%); m.p. 185-187 °C; IR (v_{max}/cm^{-1}) 3056 (C-H aromatic), 1649 (C=N) and at 1598 (C=C); ¹HNMR (400 MHz, DMSO-d₆) δ *ppm*: 8.06 (d, *J* = 9.3 Hz, 1H, Ar-H), 7.99 (d, *J* = 7.6 Hz, 2H, Ar-H), 7.83 (d, *J* = 9.3 Hz, 1H, Ar-H), 7.68 (d, *J* = 7.6 Hz, 2H, Ar-H), 7.33-7.45 (m, 5H, Ar-H), 7.21-7.26 (m, 2H, Ar-H), 7.03-7.09 (m, 2H, Ar-H), 5.92 (dd, *J* = 5.5, 12.2 Hz, 1H, pyrazoline-CH), 3.98 (dd, *J* = 12.2, 17.9 Hz, 1H, one proton of pyrazoline-CH₂), 3.82 (s, 3H, OCH₃), 3.31 (dd, *J* = 5.5, 17.9 Hz, 1H, one proton of pyrazoline-CH₂); MS (m/z (R.I. %)): [M]⁺ 424 (2), 421 (23.1), 135 (56.1), 57 (100); Anal. Calcd. for C₂₆H₂₁FN₄O (424.17); % C, 73.57; H, 4.99; N, 13.20, Found: % C, 73.65; H, 4.89; N, 13.31.

3.1.1.12. 3-(4,5-dihydro-3-(3-methoxyphenyl)-5-m-tolylpyrazol-1-yl)-6-phenylpyridazine (IXI).

Yellow powder (yield 66%); m.p. 195-197 °C; IR (v_{max}/cm^{-1}) 3057 (C-H aromatic), 1647 (C=N) and at 1598 (C=C); ¹HNMR (400 MHz, DMSO-d₆) δ *ppm*: 8.06 (d, *J* = 8.8 Hz, 1H, Ar-H), 7.97-8.01 (m, 2H, Ar-H), 7.84 (d, *J* = 9.3 Hz, 1H, Ar-H), 7.33-7.49 (m, 6H, Ar-H), 6.97-7.10 (m, 5H, Ar-H), 5.92 (dd, *J* = 5.5, 12.1 Hz, 1H, pyrazoline-CH), 3.98 (dd, *J* = 12.2, 17.8 Hz, 1H, one proton of pyrazoline-CH₂), 3.82 (s, 3H, OCH₃), 3.28 (dd, *J* = 5.6, 17.8 Hz, 1H, one proton of pyrazoline-CH₂), 2.26 (s, 3H, CH₃); ¹³CNMR (100 MHz, DMSO-d₆) δ *ppm*: 164.4, 159.1, 155.0, 152.7, 144.7, 136.4, 130.2, 129.9, 129.4, 128.6, 128.3, 128.0, 127.6, 126.7, 126.1, 122.0, 121.8, 118.2, 116.8, 115.3, 114.5, 63.3, 55.3, 46.7, 21.5; MS (m/z (R.I. %)): [M]⁺ 420 (5.4), 135 (78.1), 57 (100); Anal. Calcd. for C₂₇H₂₄N₄O (420.20); % C, 77.12; H, 5.75; N, 13.32, Found: % C, 77.17; H, 5.69; N, 13.21.

3.1.1.13. 3-(5-(4-chlorophenyl)-4,5-dihydro-3-(3-methoxyphenyl)pyrazol-1-yl)-6phenylpyridazine (IXm).

Buff powder (yield 67%); m.p. 144-146 °C; IR (v_{max}/cm^{-1}) 3036 (C-H aromatic), 1695 1649 (C=N) and at 1592 (C=C); ¹HNMR (400 MHz, DMSO-d₆) δ *ppm*: 8.07 (d, J = 9.3 Hz, 1H, Ar-H), 8.00 (d, J = 7.3 Hz, 2H, Ar-H), 7.85 (d, J = 9.5 Hz, 1H, Ar-H), 7.45-7.51 (m, 2H, Ar-H), 7.37-7.44 (m, 6H, Ar-H), 7.30-7.34 (m, 2H, Ar-H), 7.03 (d, J = 6.9 Hz, 1H, Ar-H), 5.93 (dd, J = 5.5, 12.10 Hz, 1H, pyrazoline-CH), 4.01 (dd, J = 12.1, 17.7 Hz, 1H, one proton of pyrazoline-CH₂), 3.83 (s, 3H, OCH₃), 3.28 (dd, J = 5.5, 18.0 Hz, 1H, one proton of pyrazoline-CH₂); ¹³CNMR (100 MHz, DMSO-d₆) δ *ppm*: 159.9, 155.8, 152.6, 152.2, 142.1, 136.8, 133.2, 132.1, 130.4, 129.3, 129.2, 128.2, 127.5, 126.2, 119.4, 116.3, 115.3, 111.7, 61.6, 55.7, 43.0; MS (m/z (R.I. %)): [M]⁺ 440 (25.8), [M+2]⁺ 442 (9.5), 306 (100), 308 (36.9); Anal. Calcd. for C₂₆H₂₁ClN₄O (440.14); % C, 70.82; H, 4.80; N, 12.71, Found: % C, 70.73; H, 4.89; N, 12.65.

3.1.1.14. 3-(5-(4-fluoro-3-methoxyphenyl)-4,5-dihydro-3-(3-methoxyphenyl)pyrazol-1-yl)-6-phenylpyridazine (IXn)

Yellow powder (yield 61%); m.p. 175-177 °C; IR (v_{max} /cm⁻¹) 3056 (C-H aromatic), 1650 (C=N) and at 1598 (C=C); ¹HNMR (400 MHz, DMSO-d₆) δ *ppm*: 8.07 (d, J = 9.3 Hz, 1H, Ar-H), 8.00 (d, J = 7.3 Hz, 2H, Ar-H), 7.86 (d, J = 9.5 Hz, 1H, Ar-H), 7.34-7.53 (m, 6H, Ar-H), 7.07-7.20 (m, 2H, Ar-H), 7.03 (d, J = 6.6 Hz, 1H, Ar-H), 6.76 (br. s, 1H, Ar-H), 5.91 (dd, J = 5.5, 11.9 Hz, 1H, pyrazoline-CH), 4.00 (dd, J = 12.2, 17.9 Hz, 1H, one proton of pyrazoline-CH₂); ¹³CNMR (101 MHz, DMSO-d₆) δ *ppm*: 159.9, 156.0, 152.6, 152.2, 149.8, 147.7, 139.9, 136.8, 133.3, 130.3, 129.3, 126.2, 126.1, 119.3, 117.9, 117.8, 116.5, 116.2, 115.4, 112.0, 111.7, 61.9, 56.4, 55.7, 43.2; MS (m/z (R.I. %)): [M]⁺ 454 (25.8), 320 (100), 170 (13.5), 115 (25.5); Anal. Calcd. for C₂₇H₂₃FN₄O₂ (454.18); % C, 71.35; H, 5.10; N, 12.33, Found: % C, 71.42; H, 4.98; N, 12.41.

3.2. In vitro cytotoxicity

In vitro cytotoxicity is made according to reported method[62]. The detailed procedures were supplied in Supplementary data.

3.3. EGFR inhibition assay:

HTRF (homogeneous time-resolved fluorescence) assay method was used to conduct inhibitory action of all newly synthesized compounds against EGFR [63]. EGFR kinase (BioAssay Systems - Kinase Assay Kit (EKIN-400)–(CA 94545, USA) and its substrate were incubated for 5 min in a buffer solution with newly synthesized compounds/Erlotinib to initiate the enzymatic reaction, then ATP was added to the reaction mixture. Hold under 30 min at room temperature. The reaction was stopped by

adding detection reagents containing EDTA for 1 h then the IC_{50} values have been determined.

3.4. Cell cycle analysis

UO-31 cells were seeded for 24 h at density of 1 x 10^5 cells per well and into six well plates. Added fetal bovine serum (FBS, 10 percent), after cells incubated at 37 °C and 5 percent CO₂. The medium was replaced with (DMSO 0.1% v/v) containing the 10 μ M of compounds IXn and IXg , then incubated for 48 h, washed with cold phosphate buffered saline (PBS), fixed with 70% ethanol, rinsed with PBS then stained with the DNA fuorochrome PI, kept for 15 min. at 37 °C. A FACS Caliber flow cytometer was then used to test samples [64].

3.5. Annexin V-FITC apoptosis assay

Cells were incubated with the tested compounds IXn and IXg for 24. Then, trypsinized and washed with cold phosphate-buffer saline (PBS), then stained with Annexin V-FITC and PI in binding buffer for 15 min. at room temperature in the dark. The samples were analyzed using the flow cytometer [65].

3.6. In vitro EIISA immunoassay for cell death modulators

The levels of anti-apoptotic marker Bcl2 as well as the apoptotic marker Bax were measured in UO-31cells treated with the molecules **IXg**, **IXn** and PG for 24 h using ELISA colorimetric kits. All the procedures were performed according to the manufacturer's instructions [66]. On the other hand, the activation of caspase 3 was measured using DRG Caspase-3 (human) ELISA (EIA-4860) kit (DRG International Inc., USA), according to the manufacturer instructions.

3.7. Molecular docking study

The crystal structures of EGFR domain co-crystallized with Erlotinib (PDB ID: 1M17) were retrieved from the RCSB Protein Data Bank [67, 68]. The bound water and ligands not active in binding have been eliminated from the protein and added polar hydrogen. Then, the protein was prepared in MOE with default options for docking study using Protonate 3D protocol. The entire EGFR complex had been defined as a docking receptor binding site. Docking was done using Triangle Matcher placement

method and London dG scoring function. The docking setup was initially validated by redocking the co-crystallized ligand (Erlotinib) in the vicinity of the enzyme binding site, then removing the Erlotinib molecule and inserting the target compound during the molecular docking process. Types of docked protein interactions with ligand were analyzed after molecular docking ended.

4. Conclusion

New series of pyridazine hybridized with pyrazoline derivatives were designed and synthesized based on targeting EGFR. Preliminarily screening results at NCI, USA showed the sensitivity of Ovarian cancer IGROV1, Breast cancer T-47D, Renal cancer CAKI-1 and UO-31 cell lines to almost all screened compounds. EGFR inhibition for the promising cytotoxic compounds showed that compounds **IXn**, **IXg**, **IXb** and **IXI** exhibited inhibition effect more than Erlotinib. Also, compounds **IXg** and **IXn** showed high cell accumulation at phase G2/M phase in UO-31cells confirming its arresting cell cycle at G2/M phase. Furthermore, compounds **IXg** and **IXn** elevated Bax/Bcl2 ratio and show a pronounced enhancement in the levels of the active caspases -3 which confirmed the mechanistic pathway of them. Molecular docking study of compounds **IXg** and **IXn** showed high degrees of superimposition Erlotinib into EGFR active sites. These results were in a perfect match with the EGFR enzyme inhibition assay.

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Supplementary data.

Supplementary materials associated with this research including NCI cytotoxicity results and elemental and spectroscopic analyses are available.

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Declaration of Competing Interest:

The authors declare no conflict of interest.