DESIGN AND SYNTHESIS OF PHTHALAZINE BASED DERIVATIVES AS POTENTIAL ANTICANCER AGENTS

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Abstract

In an attempt to develop novel targeted anticancer agents, a series of novel substituted phthalazine derivatives has been designed and synthesized as inhibitors of vascular endothelial growth factor receptor (VEGFR) kinase enzyme in accordance to SAR studies of known VEGFR inhibitors. The designed compounds incorporated a biarylamide or biarylurea tail linked to a phthalazine scaffold via an amino or ether linkage or else incorporated an N-substituted piperazine motif at position 1 of the phthalazine core. The prepared compounds were evaluated for their enzymatic inhibition of VEGFR-2 kinase. Furthermore, three of the phthalazines bearing a biarylurea (**6b**, **6e &7b**) exhibited excellent broad spectrum cell growth inhibition against NCI full 60 cell panel with GI_{50} values between 0.15-5µM. In addition, docking studies were performed through docking of the investigated compounds into VEGFR-2 binding site in order to gain further insight into their binding affinities and binding interactions with the VEGFR-2 kinase.

Key words: Substituted phthalazines, Cytotoxic activity, Kinase inhibitors, VEGFR-2

Introduction

Cancer is one of the major leading causes of death worldwide(ShulanZhang, 2010). It is estimated that cancer accounts for one of every five deaths. Moreover, it is estimated that the annual number of deaths due to cancers will increase from 7.6 million in 2008 to 13 million in 2030(Ales Imramovský 2013). Despite improved imaging and molecular diagnostic techniques, and advances in prevention and chemotherapeutic management, the disease still affects many millions of patients worldwide (Nakhjiri et al., 2012). Apart from surgical treatment and irradiation techniques, chemotherapy still remains an important option for cancer therapy.

It was suggested that the vast catalog of cancer cell genotypes are manifestation of six essential alterations in cell physiology that collectively dictate malignant growth; self-sufficiency in growth signals, insensitivity to growth-inhibitory (antigrowth) signals, evasion of programmed cell death (apoptosis), limitless replicative potential, sustained angiogenesis, and tissue invasion and metastasis. Each of these physiologic changes acquired during tumor development can represent a successful breaching of an anticancer defense mechanism hardwired into cells and tissues(Hanahan& Weinberg, 2011).

Vascular endothelial growth factor (VEGF) has been considered to play a major role in angiogenesis, the process of sprouting of new blood vessel formations from preexisting ones. This process begins with the dilatation of pre-existing capillaries and venules, followed by an increase of vascular permeability. Endothelial cells, which constitute the vessels, become activated, detach, migrate, and proliferate to form new sprouts(Zhang et al., 2009,Qun Zhao, 2010,Musumeci et al., 2012). Normal angiogenesis occurs during ovulation, wound healing, and reproduction. Physiological angiogenesis is tightly controlled by a complex balance among endogenous proangiogenic and antiangiogenic factors(Clarke& Hurwitz, 2013). Aberrant angiogenesis is present in a wide range of diseases including retinopathies, arthritis, endometriosis, atherosclerosis, and cancer (Potente, 2011).

One approach for inhibiting or reducing angiogenesis involves retarding, modulating or inhibiting the effects of VEGF binding at the VEGF receptor 2 (VEGFR-2); also known as the kinase insert domain receptor or (KDR), a receptor tyrosine kinase expressed on vascular endothelial cells (Francesca Musumeci, 2012).

Binding of VEGFs to their specific VEGFRs induces receptor dimerization, resulting in modification in the intracellular domain conformation. These conformational changes lead to the exposure of an ATP-binding site, followed by ATP binding and auto- or transphosphorylation on specific tyrosine residues on the receptor dimmers (Holmes, 2007) and on downstream signal transduction proteins (Koch, 2011). This mechanism leads to the initiation of a typical receptor signal transduction cascade, which activates several downstream enzymatic pathways, including p38MAPK, Raf/MEK/ERK, and PI3K/PKB pathways(Tomao et al., 2014).Thus inhibition of VEGFR kinase has been considered as an attractive target in developing novel anticancer therapeutics.

Multiple reports have detailed several small-molecule inhibitors of VEGFR-2 acting by binding to the ATP binding site in the intracellular kinase domain resulting in diminished VEGF signal transduction, these inhibitors may be broadly categorized into two main types; type I inhibitors represent ATP competitors that generally bind in or around the catalytic site of the kinase in its active conformation, in the region originally occupied by the adenine moiety of ATP (Zhang et al., 2009). On the other hand, type II inhibitors stabilize the inactive conformation of the enzyme derived upon the movement of the DFG motif, hence, they exploit new interactions with the lipophilic pocket revealed in this new rearrangement(Jabbour et al., 2013).

In the past few years, a number of VEGFR-2 inhibitors have proven clinical success. Among these, Vatalanib (PTK-787) (I) (Fig. 1), the 1,4-disubstituted phthalazine inhibitor, which is in Phase III clinical trials for metastatic colorectal cancer (Scott et al., 2007).

The biarylurea based inhibitor sorafenib (II)(Fig. 1), has been approved for treatment of advanced renal cell carcinoma (Brian B. Hasinoff 2010). Interestingly, sorafenib is not a specific inhibitor of its primary target enzyme, but also displays potent inhibitory activity for the kinase domain of the B-raf and platelet derived growth factor receptor beta (PDGFRB), which may also contribute to its antiangiogenic effects. The

promising clinical efficacy of sorafenib may be attributed to the combination of VEGFR1-2, PDGFR- β , and B-raf (Ott et al., 2010).

Interest in developing novel biarylurea-based VEGFR-2 inhibitors has been increasingly highlighted with the clinical success of other sorafenib analogues as regorafenib(**III**) and linifanib (**IV**)(Chuma et al., 2014). Also, the quinazoline derivative (**V**) was reported as a potent inhibitor of VEGFR-2 in addition to being efficacious in HepG2 human tumor xenograft model in BALB/c-nu mice(Yu et al., 2012).

Moreover, AAL993 (VI)is known as a hybrid-design type II inhibitor derived from the type I inhibitor vatalanib. It is considered as a potent inhibitor of VEGFR family that was identified after main optimization of an anthranilamide series(Blanc et al., 2013).

From another point of view, the piperazinyl moiety, the small rigid heterocyclic motif, has been successfully incorporated in several potent kinase inhibitors(L. Garuti, 2009,Shallal& Russu, 2011,Rice et al., 2012,Zhang et al., 2010,Pollard& Mortimore, 2009).The aryl piperazine based derivative (VII) (Fig. 1), has been reported to exhibit potent single digit nanomolar inhibitory activity on specific kinases(Musumeci& Radi& Brullo& Schenone, 2012,Abouzid et al., 2012).In a recent disclosure, the SAR and synthesis of pyrazolopyrimidines as potent and selective AKT1 and p70S6K kinase inhibitors, of which compound (VIII) also demonstrated potent *in vivo* tumor growth inhibition(Rice& Kim& Bussenius& Anand& Blazey& Bowles& Canne-Bannen& Chan& Chen& Co& Costanzo& DeFina& Dubenko& Engst& Franzini& Huang& Jammalamadaka& Khoury& Klein& Laird& Le& Mac& Matthews& Markby& Miller& Nuss& Parks& Tsang& Tsuhako& Wang& Xu, 2012).



Fig. 1: Structures of some potent tyrosine kinase inhibitors.

1. Rationale and design

Based on the above findings, three series of phthalazine-based derivatives linked to abiarylamide or biarylurea tail to position 1 of the phthalazine core via an amino or ether linkage ehave been designed and synthesized as VEGFR-2 kinase inhibitors. In addition, different substitution pattern have been introduced to the terminal aromatic ring aiming to better occupy to the hydrophobic pocket revealed by the kinase in its DFG-out conformation, (**Fig 2**).The design of these series was based on the study of structure activity relationships as well as common pharmacophoric features shared by several type II VEGFR inhibitors which involve; a flat heteroaromatic ring system that occupies the ATP binding region of the kinase, hydrogen bond donor-acceptor pair represented by the urea or amide moieties which usually form hydrogen bonds with Glu 885 and Asp 1046 residues, in addition to a substituted terminal aryl moiety to occupy the hydrophobic pocket revealed by the kinase in its

inactive conformation (D.M. Sammond, 2005). Furthermore, a fourth series of substituted piperazine based derivatives was also designed to target the human kinome via linking the two major fragments; the phthalazine core, recognized as a kinase privileged fragment, to the piperazine moiety aiming to offer morerigidity and to facilitate deriving spatial-activity relationships. Thus, phthalazine based derivatives bearing substituted piperazinyl moieties were also designed (Shallal& Russu, 2011).



Fig. 2: Strategy for the design of the target compounds 5a-d, 6a-f and 7a-f.

3. Results and discussion

3.1. Chemistry

The pathways adopted for the synthesis of the new substituted phthalazine derivatives are depicted in Schemes(1a,b). The key intermediates 1-aryl-3-(4hydroxyphenyl)urea (1a-c), were prepared following the literature methods as illustrated in Scheme 1a(Stout et al., 1985). On the other hand cyclization of the respective ortho substituted benzoic acid derivative with hydrazine hydrate afforded the corresponding phthalazinones (2a,b). Chlorination of 2a,b with phosphorous oxychloride afforded the corresponding1-chlorophthalazines(3a,b) which were used to prepare the target series of 1-substituted phthalazines via reaction with various nucleophiles. Thus, the reaction of **3a,b** with p-phenylenediamine in butanol afforded the N^{1} -(phthalazin-1-yl)benzene-1,4diamines (4a,b). The latter compounds (4a,b) were further reacted with the respective benzoyl chloride in acetonitrile, in presence of triethylamine to afford the target phthalazines bearing the biarylamide tail(**5a-d**). Moreover, the reaction of (**4a,b**) with different isocyanates in DMF yielded the biarylurea(6a-f). On the other hand, 1-Aryl-3-(4-(4-substituted phthalazin-1-yloxy)phenyl)urea(7a-f) were obtained via reacting the respective 1-chlorophthalazines (3a,b) with the appropriate intermediates (1a-c) in refluxing acetonitrile. Finally, refluxing the1-chlorophthalazines(3a,b) with the appropriate piperazine derivative in ethanol yielded the target phthalazines bearing the substituted piperazinyl tail (**8a-j**)in a good yield (**Scheme 1b**).



Scheme 1a: Reagents and conditions: i) dioxane, rt, 1h.



Scheme 1b: Reagents and conditions: i) NH_2NH_2 99%, propanol, w.b., 2h ii)POCl₃, reflux 70 °C, 2h iii) p-phenylenediamine, butanol, w.b., 1h, iv) benzoyl chlorides, acetonitile, triethylamine v)phenylisocyanates, DMF, reflux, 8h, vi) **1a-c**, Cs₂CO₃, acetonitrile, reflux, 6h, vii) piperazines, K₂CO₃, KI, ethanol, reflux, 3h.

3.2. Biological Evaluation

3.2.1. In vitro anticancer screening at full NCI 60 cell panel

Aiming to assess the general antitumor activity of the synthesized phthalazines, the structures of the final compounds were submitted to National Cancer Institute "NCI" (<u>www.dtp.nci</u>. nih.gov), Bethesda, Maryland, USA. *In vitro* NCI anticancer screening is a two-stage process, beginning with the evaluation of the selected compounds against the full NCI 60 cell lines panel representing leukemia, Non-Small Cell Lung Cancer, melanoma, colon cancer, CNS cancer, breast cancer, ovarian cancer, renal cancer and prostate cancer at a single dose of 10 μ M. NCI selection is based on degree of structure variation and application of computer modeling techniques to prioritize compounds based on their ability to add diversity to the NCI small molecule compound collection.

Twelve of the submitted phthalazines (**5b**, **5d**, **6b**, **6e**, **7b**, **7e**, **8a**, **8d**, **8f**, **8g**, **8h and 8i**) were selected under the NCI drug discovery program for screening against the full NCI 60 cell panel at single dose 10μ M. Among the investigated twelve compounds, three compounds (**6b**, **6e and 7b**) were selected by NCI for further screening at 5-log dose molar range due to their selective prominent cell growth inhibition on leukemia, renal and breast cell lines.

3.2.2.1. Primary in vitro antineoplastic single dose assay.

Primary in vitro single dose anticancer assay was performed in full NCI 60 cell panel. Results for each compound were reported as a mean graph of the percent growth of the treated cells compared to the untreated control cells. Three of the investigated substituted phthalazine-based derivatives showed a distinctive pattern of sensitivity against different NCI cell lines(Table 1). The biarylurea based derivative (6b) linked to the phthalazine core via an NH linker exhibited remarkable broad spectrum cell growth inhibition above 90% against various cell lines including leukemia, non-small cell lung cancer, melanoma, prostate cancer and breast cancer cell lines. In addition, its analogue (6e) having a methyl substituent at 4-position of phthalazine scaffold exhibited even more potent and broad spectrum inhibition above 100% against a wide variety of NCI cell lines including leukemia, colon cancer, CNS cancer, melanoma, renal cancer and breast cancer cell lines. Interestingly, it showed highly potent inhibition above 100% against most of the tested leukemia and melanoma cell lines. Furthermore, the biarylurea analogue (7b) linked to the phthalazine nucleus via an ether linkage also showed excellent mean growth inhibition of 106.99% due to its broad spectrum inhibition against most of the tested NCI cell lines representing all the nine tumor subpanels specially those of leukemia, colon, melanoma and breast cancer cell lines. On the contrary, compounds 8a, 8d, 8f, 8g, 8h, 8i have not shown any significant cell line growth inhibition under the same test conditions.

Table 1: *In vitro* NCI 60 cell line anticancer screening results of compounds (**5b**, **5d**, **6b**, **6e**, **7b**, **7e**, **8a**, **8d**, **8f**, **8g**, **8h**, **8i**) at single dose of 10 μ M presented as percent cell growth promotion.

Panel/Cell Line	5b	5d	6b	6e	7b	7e	8a	8d	8f	8g	8h	8i
Leukemia												
CCRF-CEM	103.12	90.95	13.13	11.02	8.70	40.21	96.87	102.23	102.72	103.32	99.65	102.09
HL-60(TB)	102.89	102.77	18.13	-34.81	3.65	80.06	107.55	96.63	113.97	103.11	110.23	108.28
K-562	102.10	100.79	9.49	-16.67	5.58	61.10	110.30	106.14	100.18	115.93	110.26	101.01
MOLT-4	97.31	83.48	-1.61	9.76	3.99	36.08	98.63	106.18	108.68	110.50	102.08	96.07
RPMI-8226	104.44	89.35	8.62	-11.88	-0.08	55.63	100.96	107.04	96.92	104.39	100.71	103.82
SR	94.95	86.34	17.49	-3.51	2.64	38.47	102.08	102.81	100.59	107.29	95.10	92.08
Non-Small Cell Lung Cancer												
A549/ATCC	100.66	97.94	20.05	26.37	40.00	61.53	99.63	100.43	94.07	99.29	88.97	92.29
HOP-62	91.79	88.80	20.57	44.24	26.74	71.07	107.44	110.27	100.46	97.06	100.11	99.80
HOP-92	97.54	55.36	38.66	4.62	12.92	93.48	103.83	102.24	112.38	95.91	112.12	119.07
NCI-H226	77.91	76.26	33.32	71.92	-54.51	79.11	100.15	98.08	102.75	95.24	90.14	98.72
NCI-H23	98.46	95.03	12.46	48.79	0.50	84.87	104.58	104.78	104.13	102.53	100.60	105.95
NCI-H322M	109.04	111.61	64.33	47.83	60.93	82.41	113.25	113.62	100.59	112.45	120.92	107.42
NCI-H460	101.43	96.41	9.89	26.11	21.50	78.22	104.65	107.58	104.04	105.33	101.10	103.32
NCI-H522	96.77	94.73	11.92	30.26	-46.35	63.79	99.80	99.42	93.81	94.50	88.96	94.54
Colon Cancer												
COLO 205	104.12	90.33	18.58	28.17	11.92	87.36	109.27	106.06	108.94	97.72	120.10	113.40
HCC-2998	113.44	125.00	30.69	40.46	-42.18	109.78	116.92	117.02	99.78	108.63	100.99	101.69
HCT-116	100.08	92.15	15.20	7.67	-56.86	72.47	99.11	102.65	103.09	112.65	95.64	98.11
HCT-15	108.95	101.25	18.27	6.39	2.76	69.54	110.84	107.84	98.61	98.47	91.84	90.29
HT29	104.37	94.32	32.43	2.24	0.52	76.99	105.50	98.23	95.98	98.30	90.03	91.22
KM12	102.76	101.06	18.93	14.10	-32.31	57.68	104.00	104.10	107.39	108.60	113.05	110.46
SW-620	102.08	92.70	26.99	20.31	-30.09	95.03	102.08	103.69	103.84	106.80	106.38	112.75
CNS Cancer												
SF-268	98.05	90.01	42.42	48.73	12.64	70.55	104.16	102.79	101.06	107.58	110.27	105.88
SF-295	101.82	104.68	17.82	4.80	40.69	91.83	102.13	106.90	94.17	96.02	92.70	87.83
SNB-19	104.47	92.02	32.96	16.20	-76.47	84.08	102.85	113.07	107.18	104.97	99.86	100.77
SNB-75	98.06	81.20	46.25	68.13	-3.13	47.27	104.44	104.58	101.30	98.14	108.60	113.97
U251	99.49	81.02	18.31	5.40	-31.11	60.02	99.27	96.94	96.04	96.73	96.20	98.57
Melanoma												
LOX IMVI	99.86	91.92	33.64	-27.27	-77.27	78.14	98.66	97.47	106.33	102.67	105.88	102.07
MALME-3M	99.61	98.95	38.46	-9.90	0.78	92.60	109.49	109.37	102.02	114.03	103.06	107.11
M14	99.68	97.11	19.31	37.26	7.30	79.40	110.29	102.33	104.47	94.84	95.05	98.56
MDA-MB-435	102.80	101.78	13.74	-39.98	-33.34	79.17	104.97	104.25	97.40	99.52	93.54	99.61
SK-MEL-28	114.13	106.32	44.13	-11.92	-31.58	88.26	115.55	108.44	113.30	108.85	104.25	101.01
SK-MEL-5	99.78	78.70	7.12	52.34	-43.28	70.24	102.99	97.51	97.52	98.10	92.88	97.04
UACC-257	92.99	97.43	15.33	73.78	34.44	83.89	95.84	99.00	92.41	92.96	95.34	98.51
UACC-62	89.82	95.86	13.41	-34.42	-14.30	45.80	102.94	99.57	109.16	109.52	99.19	101.37

Ovarian Cancor												
	101.65	101.02	39.69	61.98	8.16	90.91	119.57	116.24	108.99	112.09	108.84	112.06
IGROVI	100.69	93.28	17.58	44.40	-62.82	69.82	104.89	99.94	112.08	108.49	109.78	114.42
OVCAR-3	105.82	105.30	11.14	42.61	-20.81	70.77	106.79	109.27	97.80	103.54	91.81	100.46
OVCAR-4	11234	98.51	85.59	72.85	-16.00	119.12	128.13	105.20	95.04	98.21	87.87	98.16
OVCAR-5	96.15	93.11	18.24	31.43	19.42	76.04	97.65	97.79	102.44	97.13	95.29	98.86
OVCAR-8	102.88	96.02	36.17	45.08	37.64	72.79	102.05	108.96	108.60	107.36	96.87	104.78
NCI/ADR- RES	87.34	83.46	38.21	66.42	56.69	69.51	104.75	103.94	101.30	96.47	104.74	108.00
SK-OV-3												
Renal Cancer												
786-0	106.81	109.40	50.39	-27.85	N/A	83.90	106.39	103.55	93.29	100.35	90.25	92.37
A498	98.67	88.94	24.54	35.41	24.52	51.51	105.00	97.78	76.51	71.73	65.35	N/A
ACHN	111.96	111.40	22.43	58.77	-15.09	80.72	114.90	115.42	106.81	100.38	98.42	99.63
CAKI-1	94.57	86.32	20.58	33.85	18.67	79.66	96.10	108.56	90.31	95.26	86.85	94.80
RXF 393	115.04	98.74	65.65	-39.72	-91.21	81.98	101.64	108.23	107.83	104.83	99.82	98.58
SN12C	100.36	90.98	26.41	60.11	-3.83	79.10	109.76	102.35	100.92	100.48	100.40	104.22
TK-10	116.14	109.44	31.96	51.66	27.53	84.79	109.81	104.97	90.61	97.25	88.69	83.19
UO-31	78.63	57.84	18.97	51.21	-51.60	51.36	91.35	96.00	92.23	107.89	95.92	102.32
Prostate Cancer												
PC-3	91.56	72.64	1.46	22.81	-29.02	38.97	92.56	101.38	100.95	99.43	98.99	100.32
DU-145	113.58	101.68	33.71	57.80	11.15	81.39	104.50	99.03	100.32	111.19	106.90	110.75
Breast Cancer												
MCF7	92.95	84.53	5.46	24.87	14.27	72.69	102.79	105.33	92.06	96.67	86.47	89.07
MDA-MB	93.31	73.03	36.57	23.06	-25.22	52.88	107.72	107.61	113.82	127.47	120.61	117.24
231/ATCC	104.82	89.72	58.32	3.43	37.47	76.42	112.75	108.80	106.23	105.12	110.06	108.69
HS 578T	106.63	92.23	15.70	11.41	-10.33	71.99	111.77	103.78	66.68	75.55	N/A	N/A
BT-549	97.57	79.82	-6.32	8.43	10.20	57.90	99.99	101.10	87.22	90.54	82.33	95.97
T-47D	101.43	103.76	0.17	6.44	-76.16	68.60	97.68	93.33	103.56	98.82	84.75	87.37
MDA-MBA- 468												
Mean	100.80	93.03	25.03	23.23	-6.99	72.80	104.90	104.15	100.54	101.90	99.05	101.38

* Results are presented as percent cell growth promotion.

* Values less than 10% growth promotion are presented in bold.

3.2.2.2. In vitro 5 log dose full NCI 60 cell panel assay.

The three compounds(**6b**, **6e & 7b**) were selected by NCI for further 5 log dose screening against full NCI 60 cell panel. All investigated compounds revealed potent antiproliferative activity against most of the tested cell lines representing the nine different subpanels with GI₅₀ values between "0.15-5 μ M". All the 60 cell lines, representing nine tumor subpanels, were incubated at five different concentrations (0.01, 0.1, 1, 10 &100 μ M) of each of the tested compounds. The outcomes were used to create log concentration vs % growth inhibition curves and three response parameters (GI₅₀, TGI and LC₅₀) were calculated for each cell line, (**Table 2**). The biarylurea-based derivative (**6b**) linked to the phthalazine core via an NH linker exhibited remarkable GI₅₀ between "0.82-5 μ M" against the investigated cell panels with submicromolar GI₅₀values of 0.82 & 0.95 μ M against the melanoma LOX IMVI &UACC-62 cell lines, respectively. Moreover, its analogue (**6e**) showed potent single digit micromolar GI₅₀values between "1.61-4.77 μ M"against all the tested cell lines

With regard to the sensitivity against some individual cell lines, compound (**7b**) exhibited the highest broad spectrum submicromolar inhibitory activity against most of the tested cell lines specially those of leukemia, colon, melanoma and breast cancer cell panels with GI_{50} values (0.15-2.81) (**Table 2**).

As for the selectivity of the test compounds towards some specific tumor subpanels which is calculated based on the ratio obtained by dividing the full panel MID (the average sensitivity of all cell lines toward the test agent) by their individual subpanel MID (the average sensitivity of all cell lines of a particular subpanel toward the test agent) (M.C. Alley, 1988,M.R. Grever, 1992). As per this criterion, the test compounds (**6b**, **6e** &**7b**) were regarded to be more selective against leukemia, renal, melanoma and breast cancer subpanel. It was reported that levels of VEGFR are over expressed in cell lines derived from leukemia, melanoma and renal cancer (Min Sun, 2010). Thus, the cytotoxicity patterns of these compounds may support the original VEGFR targeting strategy.

Panel Cell Line		Compound 6b			Compound 6e			Compound 7b		
		GI ₅₀	TGI	LC ₅₀	GI ₅₀	TGI	LC ₅₀	GI ₅₀	TGI	LC ₅₀
Leukemia	CCRF-CEM	2.57	17.0	>100	2.38	5.99	>100	0.32	ND	>100
	HL-60(TB)	2.42	7.00	>100	2.03	4.18	8.64	1.08	13.4	>100
	K-562	1.68	26.4	>100	2.20	4.67	9.92	0.37	>100	>100
	MOLT-4	2.09	6.36	>100	2.36	6.00	>100	0.37	>100	>100
	RPMI-8226	2.50	8.75	>100	2.60	6.48	>100	0.48	4.58	>100
	SR	3.89	23.3	>100	2.56	9.55	>100	0.40	>100	>100
Non-Small	A549/ATCC	3.03	19.3	>100	3.24	8.59	37.9	2.81	11.5	>100
Cell Lung Cancer	EKVX	2.95	25.3	>100	3.47	17.9	87.8	1.61	4.50	>100
	HOP-62	3.00	14.9	74.9	1.81	3.70	7.56	1.09	4.32	47.7
	НОР-92	2.55	7.26	>100	2.56	6.60	42.0	1.64	4.37	17.1

Table 2: *In vitro* NCI 5 log dose results for compounds (**6b**, **6e & 7b**) in µM against 60 cell panel.

	NCI-H226	2.46	12.5	43.2	3.61	15.9	64.6	0.62	5.42	>100
	NCI-H23	3.07	14.4	71.8	2.38	6.31	40.9	0.75	2.87	9.61
	NCI-H322M	5.00	15.0	>100	3.31	1.25	69.5	2.60	9.51	34.9
	NCI-H460	2.91	11.9	56.8	3.46	11.7	39.1	2.79	8.32	45.5
	NCI-H522	1.32	8.81	43.2	1.84	3.88	8.21	0.19	0.41	0.91
Colon	COLO 205	2.77	9.58	34.6	2.22	5.29	19.2	1.18	2.92	7.21
Cancer	HCC-2998	3.26	19.8	>100	2.84	8.09	27.8	1.84	3.48	6.57
	HCT-116	2.46	12.0	37.4	1.86	3.59	6.93	0.32	1.20	4.65
	НСТ-15	1.59	11.8	71.5	1.91	4.53	13.9	0.96	3.08	9.67
	НТ29	2.65	14.1	60.0	1.78	3.67	7.56	0.46	2.39	>100
	KM12	1.47	11.4	45.0	1.99	4.53	11.1	0.88	2.65	7.38
	SW-620	3.16	18.6	81.8	2.15	4.76	11.9	0.34	1.06	30.3
CNS	SF-268	3.19	28.1	>100	2.57	7.17	47.0	1.35	7.21	70.2
Cancer	SF-295	2.84	11.0	45.8	2.04	4.21	8.68	2.54	6.27	36.9
	SF-539	2.88	11.5	36.6	1.61	3.32	6.83	0.28	1.13	3.81
	SNB-19	3.67	19.0	>100	2.77	8.72	56.6	1.74	4.79	58.3
	SNB-75	1.88	8.67	31.8	1.91	5.49	21.3	0.45	1.95	5.20
	U251	2.86	10.3	37.2	1.98	3.75	7.07	1.01	2.38	5.61
Melanoma	LOX IMVI	0.82	14.6	53.2	1.75	3.27	6.11	0.16	0.45	1.63
	MALME-3M	3.94	16.5	50.0	2.17	4.79	20.0	0.44	2.31	9.69
	M14	1.52	11.5	40.7	1.76	3.57	7.27	1.12	3.16	8.93
	MDA-MB-	2.14	9.42	30.9	1.73	3.43	6.81	0.29	1.07	5.65
	435 SV MEL 2	2.70	7.70	44.8	2.22	4.59	9.50	1.57	3.65	8.50
	SK-MEL-2	3.04	12.8	51.5	1.63	3.20	6.29	0.42	1.65	5.33
	SK-WIEL-20	2.31	6.82	24.7	1.85	3.50	6.65	1.02	2.81	6.56
	SK-WIEL-5	3.41	10.8	33.5	2.39	5.30	1.48	0.15	4.04	12.9
	UACC 62	0.95	7.97	38.1	1.75	3.59	7.35	1.08	3.46	14.0
Overien	IGROV1	2 52	30.7	>100	3 98	19.5	>100	1 16	5.00	>100
Cancer	OVCAR-3	2.52	10.3	70.0	1.87	3 99	× 100	0.35	1.15	3 95
	OVCAR-4	1.97	8 17	>100	2.79	8 36	58.2	0.55	3 49	17.0
	OVCAR-5	4 15	34.3	>100	2.10	4 99	18.7	1.89	4 47	12.6
	OVCAR-8	3.12	12.4	>100	3 35	9.32	47.4	1.10	13.6	>100
	NCI/ADR-	2.79	26.3	>100	3.63	14.9	>100	2.63	>100	>100
	RES	3.05	17.7	>100	2.61	6.59	30.7	1.74	5.48	27.2
	SK-OV-3	0100	1,11,	/ 100	2101	0.07	2011		0110	
Renal	786-0	3.09	25.2	>100	1.71	3.34	6.55	0.87	2.35	ND
Cancer	A498	3.79	18.8	>100	3.21	10.5	38.5	1.50	3.06	6.26
	ACHN	1.12	7.12	>100	1.84	3.69	7.41	1.05	2.33	5.18
	CAKI-1	1.97	12.5	88.0	3.49	12.8	43.6	2.72	8.70	43.9
	RXF 393	4.28	21.4	>100	2.17	4.13	7.84	0.31	1.05	3.74

	SN12C	4.58	37.2	>100	2.86	9.41	60.1	1.03	3.58	>100
	TK-10	5.03	24.4	>100	3.19	9.42	33.7	2.12	4.28	8.64
	UO-31	1.90	15.8	>100	2.14	6.01	31.8	1.29	2.67	ND
Prostate	PC-3	2.32	11.0	55.4	3.49	11.9	62.7	1.64	6.08	35.6
Cancer	DU-145	3.17	26.9	>100	4.77	19.5	66.9	0.73	2.08	4.88
Derret	MOET	1 42	0.22	64.1	2.21	10.0	(9.7	0.22	1.90	ND
Breast	MCF/	1.45	8.32	04.1	2.21	10.9	08.7	0.32	1.89	ND
Cancer	MDA-MB	3.40	15.9	64.9	2.38	5.94	33.9	0.39	3.68	42.4
	231/ATC C	3.83	27.7	>100	2.52	6.40	>100	0.69	47.0	>100
	HS 578T	2.15	9.28	>100	2.21	5.55	66.0	2.08	7.43	58.1
	BT-549	1.03	5.35	79.3	2.06	5.08	56.4	0.24	7.68	>100
	T-47D	2.57	9.47	>100	2.84	7.43	36.2	0.75	2.52	7.18
	MDA-MB- 468									

*The GI_{50} value (growth inhibitory activity) corresponds to the concentration of the compound causing 50% decrease in net cell growth.

*The TGI value (cytostatic activity) is the concentration of the compound resulting in total growth inhibition.

*LC₅₀ value (cytotoxic activity) is the concentration of the compound causing net 50% loss of initial cells at the end of the incubation period of 48 h.

*Submicromolar GI₅₀ values are presented in bold.

3.2.2. In vitro VEGFR tyrosine kinase activity

The VEGFR tyrosine kinase assays were performed at BPS Bioscience (www.bpsbioscience.com). All the synthesized compounds representing the four series of phthalazine based derivatives were evaluated for their ability to inhibit VEGFR-2 tyrosine kinase at single dose of 10 μ M. At this concentration, the biarylurea based derivative (**6c**) incorporating a 3-chloro-4-triflouromethyl substituent on the terminal phenyl ring has demonstrated a significant inhibition of 70% for the VEGFR kinase activity, however its analogue **7c** exhibited moderate VEGFR inhibition percent 0f 47%. The mean percent inhibition of the compounds on VEGFR at 10 μ M concentration are presented in **Table 3**.

Investigating the results of VEGFR inhibitory activity among the newly synthesized phthalazines, it was revealed that derivatives bearing a substituted biarylurea motif and linked to the phthalazine nucleus via amino linker (**6a-f**)generally tended to exhibit higher inhibition percent than their ether linked analogues (**7a-f**). And within these biarylurea based derivatives, the 4-chloro-3-triflouromethyl derivatives, having a substitution pattern similar to that of sorafenib, seemed to demonstrate the highest VEGFR inhibition percent. Unfortunately, neither of the amide based phthalazines nor the piperazine based ones exhibited any significant VEGFR inhibition.

Furthermore, compound (**6c**)which exhibited 70% VEGFR kinase inhibition was tested at 5 dose concentration to calculate its IC_{50} . Its profiling data against VEGFR2 showed increased inhibition of VEGFR activity with increasing concentration with IC_{50} = 13.4 μ M.

Compound	% inhibition	Compound	% inhibition
5a	4	7d	10
5b	8	7e	14
5c	12	7f	21
5d	15	8 a	13
6a	13	8b	8
6b	20	8c	9
6с	70	8d	5
6d	15	8e	9
6e	19	8 f	14
6f	32	8g	13
7a	5	8h	9
7b	13	8i	6
7c	47	8 j	10
Staurosporine	100		

Table 1: The VEGFR inhibition percent of the synthesized phthalazines at 10 μ M concentration.

3.3. Molecular modeling studies

Molecular docking study was performed using Accelry's Discovery Studio 2.5 software CDOCKER protocol through docking of the synthesized compounds in the VEGFR-2 kinase active site (Accelry's Discovery Studio 2.5, 2010). Docking study aimed to interpret the VEGFR-2 inhibitory activity of the investigated molecules and gain further insight into their binding affinities and binding interactions with the kinase active site. The coordinates of the VEGFR2 structure were obtained from the crystal structure of VEGFR complexed with sorafenib as its inhibitor (PDB code 4ASD), which revealed the hydrogen bond interactions between the NH and CO motifs of urea moiety with the backbone of Asp1046 and the carboxylic acid moiety of Glu885, respectively, as well as a H-bond with Cys919 residue in the hinge region of the kinase active site(D.M. Sammond, 2005).

The docking algorithm was validated via redocking of the co-crystallized ligand (sorafenib) in the VEGFR-2 active site where the docking procedure was able to retrieve the co-crystallized pose with RMSD value of 2.03 A° .

Docking of the investigated compounds revealed that the substituted urea based derivatives (6c & 7c) were capable of occupying the ATP binding site of the VEGFR-2 kinase while maintaining the essential key interactions. Thus, a hydrogen bond has been observed between carbonyl group of the urea motif with the backbone NH of Asp 1046 where as the urea NH groups have been engaged in two hydrogen bond interactions

with the carboxylic acid of Glu 885. However, the substituted biarylurea based derivatives (6c & 7c)missed the key H-bond interaction with Cys919,which may explain their relatively lower activity against VEGFR-2 inhibitory activity. On the contrary, the piperazine based derivatives (8a-j) failed to form the essential H-bond interactions with the VEGFR active site which might explain their poor inhibitory activity. One of the best scored poses of the urea based compounds (6c & 7c) are presented in Fig 3 & 4.



Fig. 3. Docking of compound **6c** in the ATP binding site of VEGFR-2 kinase(PDB code 4ASD); showing key hydrogen bonding interactions (dotted lines) betweenCO and NH motifs of urea moiety with the backbone of Asp1046 and the carboxylic acid of Glu885 residues.



Fig. 4.Docking of compound **7c** in the ATP binding site of VEGFR-2 kinase (PDB code 4ASD); showing key hydrogen bonding interactions (dotted lines) between CO and NH motifs of urea moiety with the backbone of Asp1046and the carboxylic acid of Glu885 residues.

4. Conclusion

Four series of phthalazine based derivatives bearing biarylamides(5a-d), biarylureas (6a-f) and (7a-f)or a substituted piperazine moiety at position 1 of the phathalazine nucleus (8a-j) were designed, synthesized as targeted anticancer agents targeting VEGFR kinase. Twelve of the synthesized compounds were selected by NCI for antiproliferative screening against full NCI 60 cell panel at single dose 10 µM. Three of the tested phthalazine bearing a biarylurea tail (6b,6e and 7b)exhibited excellent broad spectrum cell growth inhibition between (100% to 191%) against a variety of NCI cell lines, thus were selected for further screening at 5 log doses against full NCI 60 cell panel. The latter 5 log doses screening confirmed their excellent in vitro cell growth inhibitory activity with GI₅₀ between 0.15-5µM against most of the investigated cell lines specially those of leukemia, renal, melanoma and breast cancer cell lines. Furthermore, the synthesized phthalazines were evaluated for their VEGFR-2 inhibitory activity, which revealed a significant inhibition of 70% exhibited by the 4chloro-3-triflouromethyl substituted biarylurea based derivative (6c). Molecular docking study of 6c on VEGFR-2 kinase active site revealed its ability to form the essential H bond interactions with Glu 885 and Asp1046 key residues in the VEGFR-2 active site.

5. Experimental

5.1. Chemistry

Starting materials and reagents were purchased from Sigma – Aldrichor Acros Organics. Melting points were recorded on Gallen Kamp apparatus and were uncorrected. FT-IR spectra were recorded on a Shimadzu IR 435 spectrophotometer. ¹HNMR spectra were recorded in δ scale given in ppm on a Varian 400 MHz spectrophotometer or a Varian 300 MHz spectrophotometer. Coupling patterns aredescribed as follows: s, singlet; d, doublet, dd, doubled doublet; t, triplet; m, multiplet. *J* describes a coupling constant. The coupling constants were rounded off to one decimal place. MS spectra mass were recorded on Hewlett Packard 5988 spectrometer (70 eV). Elemental analyzes were performed at the Microanalytical Center,

Al-Azhar University. Compounds **1a-c**(Stout& Matier& Barcelon-Yang& Reynolds& Brown, 1985), **2a,b**(Smolyar& Yutilov, 2008) **and 3a,b**(Neumann, 1893) were prepared following reported procedures.

5.1.1. Synthesis of N¹-(4-Arylphthalazin-1-yl)benzene-1,4-diamines (4a,b)

General procedure:

Benzene diamine (0.33 g, 3.12 mmol, 3equiv) and the respective chlorophthalazine

(2.08 mmol, 2equiv) were treated with 2-BuOH (7.5 mL) in a tube and heated to 110 °C. The reaction quickly became a solid, yellow mass. After four hours, the reaction was cooled and diluted with water. The resultant slurry was then partitioned between equal volumes of DCM and 1N NaOH (15 mL). The aqueous layer was extracted into DCM (2x15 mL). The combined organic layers were dried over anhydrous sodium sulfate, filtered, and concentrated *in vacuo*. The resulting orange solid was crystallized from (EtOAc/pet ether 1:1) to afford the title compounds (**4a,b**) as an orange crystals.

5.1.1.1. N1-(Phthalazin-1-yl)benzene-1,4-diamine (4a)

Yield(0.21 g, 91%) ;mp 284-286°C; ¹HNMR (**300** MHz, DMSO-d₆): δ 9.15 (1H, s,-NH D₂O exchangeable), 9.04 (2H, s, -NH₂ D₂O exchangeable), δ 8.95-8.92 (2H,d, *J*= 9 Hz, phthalazine), δ 8.18-8.15 (2H,d, *J*= 9 Hz, phthalazine), δ 7.97 (1H,s, phthalazine), δ 7.65-7.62 (2H,d, *J*= 9 Hz, ArH), δ 6.97-6.83 (2H,d, *J*= 9 Hz,ArH); FT-IR ($\dot{\nu}$ max, cm⁻¹): 3400 (NH₂), 3329(NH); **MS** (Mwt.: 236.11): m/z 236.00 (M+, 43.00%), 212.00 (100.00%), 118.00 (54.00%), 56.00 (70.00%); Anal. Calcd for C₁₄H₁₂N₄: C, 71.17; H, 5.12; N, 23.71; Found: C, 71.34; H, 5.17; N, 23.96

6.1.1.2. N1-(4-Methylphthalazin-1-yl)benzene-1,4-diamine (4b)

Yield (0.22 g, 88%) ;mp 240-242°C; ¹HNMR (**300** MHz, DMSO-d₆): δ 8.88 (1H, s,-NH D₂O exchangeable), 8.85 (2H, s, -NH₂ D₂O exchangeable), δ 8.13-8.08 (2H,d, *J*= 7 Hz, phthalazine), δ 7.95-7.92 (2H,d, *J*= 7 Hz, phthalazine), δ 7.46-7.43 (2H,d, *J*= 9 Hz, ArH), δ 6.83-6.76 (2H,d, *J*= 9 Hz, ArH), δ 2.48 (3H,s, CH₃); FT-IR ($\dot{\nu}$ max, cm⁻¹): 3385 (NH₂), 3321(NH); **MS** (Mwt.: 250.12): m/z 250.00 (M+, 50.00%), 249.00 (100.00%), 108.00 (60.92%), 65.00 (77.59%); Anal. Calcd for C₁₅H₁₄N₄: C, 71.98; H, 5.64; N, 22.38; Found: C, 72.09; H, 5.78; N, 22.52.

5.1.2. Synthesis of N-(4-(4-Substituted phthalazin-1-ylamino)4-substituted phenyl)benzamide (5a-d)

General procedure:.

To a stirred solution of the respective N1-(4-arylphthalazin-1-yl)benzene-1,4diamine (**4a,b**) (10.0 mmol, 1equiv) and triethylamine (1.1 mL, 10.0 mmol) in acetonitrile (20 mL), the respective benzoyl chloride (viz.; benzoyl chloride, 4chlorobenzoyl chloride) (10.0 mmol, 1equiv) was added and the mixture was heated under reflux for 6h, until the disappearance of starting material as judged by TLC (CHCl₃/CH₃OH 9.5:0.5). The reaction mixture was filtered and then the filtrate was concentrated in vacuo, to afford the crude product which was purified by flash column chromatography (EtOAc/pet ether 9:1) to afford **5a-d** as yellow crystals.

5.1.2.1. N-(4-(Phthalazin-1-ylamino)phenyl)benzamide (5a)

Yield (0.17g; 50%), mp 142-144°C; ¹HNMR(400 MHz, DMSO-d₆): δ 9.66 (1H,s,

-NH D₂O exchangeable), δ 9.12 (1H, s, -NH D₂O exchangeable), δ 8.62 (1H, s, *J*= 8 Hz, phthalazine), δ 8.11-8.08 (2H,d, *J*= 7.6 Hz,Ar-H), δ 7.98-7.96 (2H,d, *J*= 8 Hz, phthalazine), δ 7.93-7.88 (2H, m,phthalazine), δ 7.64-7.62 (1H,m,Ar-H), δ 7.62-7.60 (2H, d, *J*= 7.6 Hz, Ar-H), δ 7.59-7.57 (2H, d, *J*= 6.8 Hz, Ar-H), δ 7.50-7.48 (2H, m,Ar-H); FT-IR ($\dot{\nu}$ max, cm⁻¹): 3550, 3510 (2NH), 1649 (C=O); **MS** (Mwt.: 340.38): m/z 340.15 [M⁺, 1.58%), 138.05 (2.70%), 86.10 (100%); Anal. Calcd for C₂₁H₁₆N₄O: C, 74.10; H, 4.74; N, 16.46; Found: C, 74.34; H, 4.80; N, 16.58

5.1.2.2. 4-Chloro-N-(4-(phthalazin-1-ylamino)phenyl)benzamide (5b)

Yield (0.20g; 54%), mp 156-158°C; ¹HNMR(400 MHz, DMSO-d₆): δ 10.50 (1H, s,-NH D₂O exchangeable), δ 9.15 (1H,s, -NH D₂O exchangeable), δ 8.99 (1H, s, phthalazine), δ 8.11-8.08 (2H, d, *J*= 9 Hz,Ar-H), δ 7.94-7.92 (2H,d, *J*= 7.6 Hz, phthalazine), δ 7.91-7.90 (2H, d, *J*= 7.6 Hz phthalazine), δ 7.56-7.54 (2H,d, *J*= 9 Hz,Ar-H), δ 7.53-7.51 (2H,d, *J*= 7.5 Hz, Ar-H), δ 7.31-7.28 (2H,m,Ar-H); FT-IR ($\acute{\nu}$ max, cm⁻¹): 3355, 3310 (2NH), 1640 (C=O); **MS** (Mwt.: 374.82): m/z 375.00 (M+, 1.57%), 245.00 (2.41%), 139.05 (46.44%), 111.10 (25.30), 64.00 (100.00%); Anal. Calcd for C₂₁H₁₅ClN₄O: C, 67.29; H, 4.03; N, 14.95; Found: C, 67.52; H, 4.08; N, 15.22.

5.1.2.3. N-(4-(4-Methylphthalazin-1-ylamino)phenyl)benzamide(5c)

Yield (0.24 g; 68%), mp 283-285°C; ¹HNMR(400 MHz, DMSO-d₆): δ 10.50 (1H, s,-NH D₂O exchangeable), δ 10.40 (1H,s, -NH D₂O exchangeable), δ 8.27-8.24 (2H, d, *J*= 7.6 Hz, phthalazine), δ 8.02-8.00 (2H, m,Ar-H), δ 7.99-7.97 (2H,d, *J*= 7.6 Hz, phthalazine), δ 7.77-7.73 (1H,d, *J*= 9.2 Hz, Ar-H), δ 7.63-7.61 (2H,d, *J*= 7.2 Hz, Ar-H), δ 7.59-7.56 (2H,d, *J*= 7.6 Hz, Ar-H), δ 7.54-7.51 (2H, m,Ar-H), δ 2.91 (3H,s, CH₃); FT-IR (δ max, cm⁻¹): 3331, 3257 (2NH), 1649 (C=O); MS (Mwt.: 354.15):m/z 354.95 (M+, 1.18%), 315.95 (15.90%), 105.00 (100.00%); Anal. Calcd for C₂₂H₁₈N₄O: C, 74.56; H, 5.12; N, 15.81; Found: C, 74.78; H, 5.19; N, 16.04

5.1.2.4. 4-Chloro-N-(4-(4-methylphthalazin-1-ylamino)phenyl)benzamide (5d)

Yield (0.20g; 51%), mp 147-149°C; ¹HNMR(400 MHz, DMSO-d₆): δ 10.65 (1H, s,-NH D₂O exchangeable), δ 10.45 (1H, s,-NH D₂O exchangeable), δ 8.05-8.03 (2H, d, *J*= 8 Hz, phthalazine), δ 7.97-7.95 (2H,d, *J*= 7.6 Hz, Ar-H), δ 7.88-7.86 (2H, d, *J*= 8 Hz, phthalazine), δ 7.63-7.61 (2H, d, *J*= 7.6 Hz, Ar-H), δ 7.59-7.57 (2H,d, *J*= 7.6 Hz, Ar-H), δ 7.31-7.28 (2H, m,Ar-H), δ 2.91 (3H, s,CH₃); FT-IR (δ max, cm⁻¹): 3300, 3192 (2NH), 1678 (C=O); **MS** (Mwt.: 388.85): m/z 388.90 (M+, 7.10%), 279.90 (7.84%), 248.95 (25.65%), 139.00 (100%); Anal. Calcd for C₂₂H₁₇ClN₄O: C, 67.95; H, 4.41; N, 14.41; Found: C, 68.24; H, 4.48; N, 14.49.

5.1.3. Synthesis of 1-Aryl-3-(4- (4-substituted phthalazin-1-ylamino)phenyl)ureas (6a-f)

General procedure:

To a stirred solution of the respective N1-(4-arylphthalazin-1-yl)benzene-1,4diamine (**4a,b**) (10.0 mmol, 1equiv) in DMF (20 mL), the respective phenyl isocyanate (viz.; phenyl- isocyanate, 4-chlorophenyl isocyanate, 3-trifluromethyl-4-chlorophenyl isocyanate) (10.0 mmol, 1equiv) was added and the mixture was heated under reflux for 6h, after which TLC (CHCl₃/CH₃OH 9:1) showed no starting material. The reaction mixture was poured over ice- water, the formed precipitate was allowed to settle, then filtered off and dried to afford the crude products(**6a-f**) which was further crystallized from EtOAc.

5.1.3.1. 1-Phenyl-3-(4-(phthalazin-1-ylamino)phenyl)urea(6a)

Yield (0.10 g; 30%), mp 145- 147°C; ¹HNMR(400 MHz, DMSO-d₆): δ 9.22 (1H, s,-NH D₂O exchangeable), δ 9.16 (1H,s, -NH D₂O exchangeable), δ 9.09 (1H,s, -NH D₂O exchangeable), δ 8.74 (1H, s, phthalazine), δ 8.68-8.66 (2H, d, *J*= 8 Hz, phthalazine), δ 8.02-8.00 (2H, d, *J*= 8 Hz, phthalazine), δ 7.58-7.56 (2H,m,Ar-H), δ 7.48-7.46 (2H, d, *J*= 7.6 Hz, Ar-H), δ 7.31-7.29 (2H,d, *J*= 7.2 Hz, Ar-H), δ 7.19-7.17 (1H,m,Ar-H), δ 6.98-6.96 (2H, d, *J*= 7.2 Hz, Ar-H) FT-IR ($\dot{\nu}$ max, cm⁻¹): 3300, 3047 (3NH), 1700 (C=O); **MS** (Mwt.: 355.14): m/z 355.10 (M+, 8.88%), 289.10 (9.51%), 123.10 (15.43%), 69.00(100.00%); Anal. Calcd for C₂₁H₁₇N₅O: C, 70.97; H, 4.82; N, 19.71; Found: C, 71.21; H, 4.89; N, 19.88

5.1.3.2. 1-(4-Chlorophenyl)-3-(4-(phthalazin-1-ylamino)phenyl)urea (6b)

Yield (0.20g; 53%), mp 168-171°C; ¹HNMR(400 MHz, DMSO-d₆): δ 10.33 (1H, s,-NH D₂O exchangeable), δ 10.07 (1H, s, -NH D₂O exchangeable), δ 9.16 (1H, s, -NH D₂O exchangeable), δ 8.96 (1H, s, phthalazine), δ 8.09-8.07 (2H, d, *J*= 9.2 Hz, phthalazine), δ 7.63-7.61 (2H, d, *J*= 9.2 Hz, phthalazine), δ 7.49-7.47 (2H, d, *J*= 8.8 Hz, Ar-H), δ 7.44-7.42 (2H,d, *J*= 8 Hz, Ar-H), δ 7.33-7.31 (d, *J*= 8.8 Hz, 2H, Ar-H), δ 6.56-6.54 (2H,d, *J*= 8 Hz, Ar-H); FT-IR ($\dot{\nu}$ max, cm⁻¹): 3396, 3341 (3NH), 1672 (C=O); MS (Mwt.: 389.10): m/z 389.00 (M+, 1.51%), 281.95 (19.41%), 253.95(100.00%), 152.95 (13.00%); Anal. Calcd for C₂₁H₁₆ClN₅O: C, 64.70; H, 4.14; N, 17.96; Found: C, 64.89; H, 4.21; N, 18.17

5.1.3.3. 1-(4-Chloro-3-(trifluoromethyl)phenyl)-3-(4-(phthalazin-1-ylamino)phenyl)urea (6c)

Yield (0.22g; 49%), mp 288-291°C; ¹HNMR(400 MHz, DMSO-d₆): δ 10.08 (1H, s,-NH D₂O exchangeable), δ 9.36 (1H, s, -NH D₂O exchangeable), δ 9.11 (1H, s, -NH D₂O exchangeable), δ 8.79 (1H, s, phthalazine), δ 8.22-8.20 (2H, d, *J*= 8.8 Hz, phthalazine), δ 8.09 (1H, s, Ar-H), δ 7.97-7.95 (1H, d, *J*= 9.6 Hz, Ar-H), δ 7.81-7.79 (2H, d, *J*= 8.8 Hz, phthalazine), δ 7.69-7.67 (1H, d, *J*= 9.6 Hz, Ar-H), δ 7.59-7.50 (2H, d, *J*= 8.4 Hz, Ar-H), δ 7.39-7.37 (2H, d, *J*= 8.4 Hz, Ar-H); FT-IR (δ max, cm⁻¹): 3321, 3263, 3126 (3NH), 1649 (C=O); **MS** (Mwt.: 457.09): m/z 457.90 (M+, 1.06%), 415.90 (7.68%), 194.95(100.00%); Anal. Calcd for C₂₂H₁₅ClF₃N₅O: C, 57.71; H, 3.30; N, 15.30; Found: C, 57.93; H, 3.28; N, 15.48

5.1.3.4. 1-(4-(4-Metyhlphthalazin-1-ylamino)phenyl)3-phenylurea(6d)

Yield (0.06 g; 20%), mp 284-286°C; ¹HNMR(300 MHz, DMSO-d₆): δ 9.20(1H, s,-NH D₂O exchangeable), δ 9.14 (1H,s,-NH D₂O exchangeable), δ 9.06 (1H, s,-NH D₂O exchangeable), δ 8.93-8.91 (1H, d, *J*= 7 Hz, phthalazine), δ 8.77-8.75 (1H, d, *J*=7 Hz, phthalazine), δ 8.22-8.20 (2H, d, *J*=7 Hz, phthalazine), δ 7.85-7.83 (2H, d, *J*= 8.4 Hz,Ar-H), δ 7.25-7.23 (1H, m, Ar-H), δ 6.95-6.93 (2H, d, *J*= 7.2 Hz, Ar-H), δ 2.81 (3H, s, CH₃); FT-IR (ψ max, cm⁻¹): 3292, 3194, 3132 (3NH), 1651 (C=O); MS (Mwt.: 369.16): m/z 369.00 (M⁺, 2.99%), 212.00 (6.90%), 93.00 (100.00%);Anal. Calcd for C₂₂H₁₉N₅O: C, 71.53; H, 5.18; N, 18.96; Found: C, 71.69; H, 5.27; N, 19.21

5.1.3.5. 1-(4-Chlorophenyl)-3-(4-(4-methylphthalazin-1-ylamino)phenyl)urea (6e)

Yield (0.20g; 50%), mp 178-180°C; ¹HNMR(400 MHz, DMSO-d₆): δ 9.20(1H, s,-NH D₂O exchangeable), δ 9.17 (1H, s, -NH D₂O exchangeable), δ 9.03 (1H, s, -NH D₂O exchangeable), δ 8.18-8.16 (2H, d, *J*= 7.6 Hz, phthalazine), δ 8.09-8.05 (2H, d, *J*= 7.6 Hz, phthalazine), δ 7.86-7.84 (2H, d, *J*= 8.8 Hz, Ar-H), δ 7.73-7.71 (2H, d, *J*= 8.8 Hz, Ar-H), δ 7.49-7.47 (2H, d, *J*= 8.8 Hz, Ar-H), δ 7.33-7.30 (2H, d, *J*= 8.8 Hz, Ar-H), δ 2.80 (3H, s,CH₃); FT-IR (\acute{v} max, cm⁻¹): 3294, 3188, 3168 (3NH), 1681 (C=O); MS (Mwt.: 403.86): m/z 403.00 (M+, 1.51%), 276.95 (17.01%), 127.00 (100.00%); Anal. Calcd for C₂₂H₁₈ClN₅O: C, 65.43; H, 4.49; N, 17.34; Found: C, 65.62; H, 4.52; N, 17.52

5.1.3.6. 1-(4-Chloro-3-(trifluoromethyl)phenyl)-3-(4-(4-methylphthalazin-1-ylamino)phenyl)urea (6f)

Yield (0.30g; 63%), mp 258-260°C; ¹HNMR(400 MHz, DMSO-d₆): δ 9.43(1H, s,-NH D₂O exchangeable), δ 9.14 (1H, s, -NH D₂O exchangeable), δ 9.02 (1H, s, -NH D₂O exchangeable), δ 8.88-8.86 (2H, d, *J*= 8 Hz, phthalazine), δ 8.24-8.22 (2H, d, *J*= 8 Hz, phthalazine), δ 8.13 (1H, s,Ar-H), δ 7.97-7.95 (1H, d, *J*= 9.2 Hz, Ar-H), δ 7.88-7.86 (1H, d, *J*= 9.2 Hz,Ar-H), δ 7.63-7.61 (2H, d, *J*= 10 Hz, Ar-H), δ 6.78-6.76 (2H, d, *J*= 10 Hz, Ar-H), δ 2.80 (3H, s, CH₃); FT-IR ($\dot{\nu}$ max, cm⁻¹): 3363, 3255, 3153 (3NH), 1681(C=O); MS (Mwt.: 471.86): m/z 472.10(M+, 0.30%), 223.00 (43.56%), 195.00 (70.63%), 52.05 (100%); Anal. Calcd for C₂₃H₁₇ClF₃N₅O: C, 58.54; H, 3.63; N, 14.84; Found: C, 58.61; H, 3.61; N, 15.03

5.1.4. 1-Aryl-3-(4-(4-substituted phthalazin-1-yloxy)phenyl)ureas (7a-f)

General procedure:

To a stirred solution of the respective chlorophthalazine derivative (**3a,b**) (10.0 mmol, 1equiv), and caesium carbonate (6.5 g, 20.0 mmol, 2equiv) in acetonitrile (20 mL), the appropriate 1-aryl-3-(4-hydroxyphenyl)urea (**1a-c**) (10.0 mmol, 1equiv) was added and the mixture was heated under reflux for 6h, after which TLC (CHCl₃/CH₃OH 9:1) showed no starting material. The filtrate was evaporated *in-vacuo* to afford the crude product (**7a-f**) which was further purified by column chromatography (using gradient elution starting from EtOAc then 1% MeOH/EtOAc).

5.1.4.1. 1-Phenyl-3-(4-(phthalazin-1-yloxy)phenyl)urea (7a)

Yield (0.22 g; 64%), mp 180-182°C; ¹HNMR(300 MHz, DMSO-d₆): δ 9.57 (1H, s, -NH D₂O exchangeable), δ 9.37 (1H, s, -NH D₂O exchangeable), δ 8.36 (1H, s,phthalazine), δ 8.10-8.00 (2H, m, phthalazine), δ 7.93-7.91 (2H, d, *J*= 7 Hz, phthalazine), δ 7.91-7.90 (2H, m, Ar-H), δ 7.49-7.47 (2H, d, *J*= 7.8 Hz, Ar-H), δ 7.19-7.25 (m, 3H, Ar-H), δ 6.89-6.86 (1H, d, *J*= 7.5 Hz,Ar-H), δ 6.67-6.65 (2H, d, *J*= 7.8 Hz,Ar-H); FT-IR (ψ max, cm⁻¹): 3304, 3235 (2NH), 1641 (C=O); MS (Mwt.: 356.38): m/z 355.00 (M-1⁺, 0.11%), 146.05 (23.71%), 118.05 (8.47%), 79.95 (100.00%);Anal. Calcd for C₂₁H₁₆N₄O₂: C, 70.77; H, 4.53; N, 15.72; Found: C,71.02; H, 4.61; N, 15.89

5.1.4.2. 1-(4-Chlorophenyl)-3-(4-(phthalazin-1-yloxy)phenyl)urea (7b)

Yield (0.19 g; 50%), mp 166-168°C; ¹HNMR(400 MHz, DMSO-d₆): δ 9.72 (1H,s,

-NH D₂O exchangeable), δ 9.44 (1H,s, -NH D₂O exchangeable), δ 8.37 (1H, s, phthalazine), δ 8.28-8.26 (2H, d, *J*= 8 Hz, phthalazine), δ 8.18-8.16 (2H, d, *J*= 8 Hz, phthalazine), δ 7.58-7.56 (2H, d, *J*= 8.8 Hz, Ar-H), δ 7.56-7.53 (2H, d, *J*= 8.8 Hz, Ar-H), δ 7.36-7.34 (2H, d, *J*= 7.2 Hz, Ar-H), δ 7.14-7.12 (2H, d, *J*= 7.2 Hz, Ar-H); FT-IR ($\dot{\nu}$ max, cm⁻¹): 3278, 3161 (2NH), 1701 (C=O); **MS** (Mwt.: 390.82): m/z 390.10 (4.47%), 365.10 (24.02%), 75.00 (100.00%); Anal. Calcd for C₂₁H₁₅ClN₄O₂: C, 64.54; H, 3.87; N, 14.34; Found: C, 64.69; H, 4.01; N, 14.62

5.1.4.3. 1-(4-Chloro-3-(trifluoromethyl)phenyl)-3-(4-(phthalazin-1-yloxy)phenyl)urea (7c)

Yield (0.22 g; 49%), mp 158-160°C; ¹HNMR(400 MHz, DMSO-d₆): δ 9.91 (1H, s,-NH D₂O exchangeable), δ 9.08 (s, 1H, -NH D₂O exchangeable), δ 8.39 (s, 1H, phthalazine), δ 8.24-8.21 (d, *J*= 8 Hz, 2H, phthalazine), δ 8.17-8.15 (d, *J*= 8 Hz, 2H, phthalazine), δ 7.99 (s, 1H, Ar-H), δ 7.85-7.83 (d, *J*= 7.6 Hz, 1H, Ar-H), δ 7.76-7.74 (d, *J*= 7.6 Hz, 1H, Ar-H), δ 7.28-7.26 (d, *J*= 8.4 Hz, 2H, Ar-H), δ 6.76-6.74 (d, *J*= 8.4 Hz, 1H, Ar-H); FT-IR ($\dot{\nu}$ max, cm⁻¹): 3300, 3153 (2NH), 1679 (C=O); MS (Mwt.: 458.82): m/z 458.10 (2.63%), 271.05 (7.73%), 195.00 (100.00%); Anal. Calcd for C₂₂H₁₄ClF₃N₄O₂: C, 57.59; H, 3.08; N, 12.21; Found: C, 57.82; H, 3.06; N, 12.57

5.1.4.4. 1-(4-(4-Methylphthalazin-1-yloxy)-3-phenyl)urea (7d)

Yield (0.20 g; 54%), mp 174-176°C; ¹HNMR(400 MHz, DMSO-d₆): δ 9.44 (2H, s, -NH D₂O exchangeable), δ 8.38-8.36 (1H, d, *J*= 8 Hz, phthalazine), δ 8.31-8.29 (1H, d, *J*= 8 Hz, phthalazine), δ 8.29-8.27 (1H, d, *J*= 8 Hz, phthalazine), δ 8.19-8.17 (1H, d, *J*= 8 Hz, phthalazine), δ 7.59-7.57 (1H, d, *J*= 8.8 Hz, Ar-H), δ 7.52-7.50 (1H, d, *J*= 8.8 Hz, Ar-H), δ 7.46-7.44 (2H, m,Ar-H), δ 7.28-7.26 (2H, d, *J*= 7.6 Hz, Ar-H), δ 6.96-6.91 (1H, d, *J*= 8.8 Hz, Ar-H), δ 6.67-6.65 (2H, d, *J*= 7.6 Hz, Ar-H), δ 2.81 (3H, s,CH₃); FT-IR ($\acute{\nu}$ max, cm⁻¹): 3302, 3141 (2NH), 1643 (C=O); MS (Mwt.: 370.40): m/z

370.00[M⁺, 2.28%), 277.95 (19.33%), 251.00 (100.00%); Anal. Calcd for $C_{22}H_{18}N_4O_2$: C, 71.34; H, 4.90; N, 15.13; Found: C, 71.49; H, 4.97; N, 15.29.

5.1.4.5. 1-(4-Chlorophenyl)-3-(4-(4-methylphthalazin-1-yloxy)phenyl)urea (7e)

Yield (0.17 g; 42%), mp 169-171°C; ¹HNMR(400 MHz, DMSO-d₆): δ 10.06 (1H, s,-NH D₂O exchangeable), δ 10.05 (1H, s,-NH D₂O exchangeable), δ 8.38-8.36 (1H, d, *J*= 7.6 Hz, phthalazine), δ 8.20-8.18 (1H, d, *J*= 7.6 Hz, phthalazine), δ 8.07-8.04 (2H, d, *J*= 7.6 Hz, phthalazine), δ 7.58-7.56 (2H, d, *J*= 8.4 Hz, Ar-H), δ 7.29-7.22(2H, d, *J*= 8.4 Hz, Ar-H), δ 7.20-7.18 (2H, d, *J*= 9.2 Hz, Ar-H), δ 6.66-6.63 (2H, d, *J*= 9.2 Hz, Ar-H), δ 2.81 (3H, s,CH₃); FT-IR (ψ max, cm⁻¹): 3300, 3153 (2NH), 1643 (C=O); **MS** (Mwt.: 404.85): m/z 404.90(M+, 2.04%), 385.85 (1.29%), 267.90 (4.75), 250.95 (28.16), 127.00 (100.00%); Anal. Calcd for C₂₂H₁₇ClN₄O₂: C, 65.27; H, 4.23; N, 13.84; Found: C, 65.52; H, 4.28; N, 13.97

5.1.4.6. 1-(4-Chloro-3-(trifluoromethyl)phenyl)-3-(4-(4-metyhlphthalazin-1-yloxy)phenyl)urea (7f)

Yield (0.20 g; 42%), mp 150-152°C; ¹H NMR(400 MHz, DMSO-d₆): δ 10.31 (1H, s,-NH D₂O exchangeable), δ 10.27 (1H, s, -NH D₂O exchangeable), δ 8.37 (1H, s,Ar-H), δ 8.19-8.16 (2H, d, *J*= 8 Hz, phthalazine), δ 8.07-8.05 (1H, d, *J*= 7.6 Hz, Ar-H), δ 7.78-8.75 (2H,d, *J*= 8 Hz, phthalazine), δ 7.61-7.59 (1H, d, *J*= 7.6 Hz, Ar-H), δ 7.56-7.54 (2H, d, *J*= 8.8 Hz, Ar-H), δ 7.24-7.19(2H, d, *J*= 8.8 Hz, Ar-H), δ 2.80 (3H, s,CH₃); FT-IR (δ max, cm⁻¹): 3300, 3153 (2NH), 1679 (C=O); MS (Mwt.: 472.85): m/z 472.80 (M+, 2.07%), 391.95 (10.86), 335.85 (25.53), 250.95 (86.76%), 194.90 (100%); Anal. Calcd for C₂₃H₁₆ClF₃N₄O₂: C, 58.42; H, 3.41; N, 11.85; Found: C, 58.49; H, 3.39; N, 12.01

5.1.5. Synthesis of Ethyl 4-(4-substituted phthalazine-1-yl)piperazine-1-carboxylate &

1-Aryl-4-(arylpiperazine-1-yl)phthalazine (8a-j)

General procedure:

To a stirred mixture of the respective 1-chlorophthalazine (**3a,b**) (10.0 mmol, 1equiv), potassium carbonate (0.27g, 20.0 mmol, 2equiv), potassium iodide (0.01g, 0.01 mmol, 0.1equiv) in absolute ethanol (20 mL), the respective piperazine (viz.; ethylpiperazinecarboxylate , phenylpiperazine, 2-flourophenylpiperazine, 2-pyridylpiperazine, 2-furoylpiperazine) (10.0 mmol, 1equiv) was added and the mixture was heated under reflux for 2h, after which TLC (CHCl₃/CH₃OH 9:1) showed no starting material. The mixture was then concentrated in vacuo, the residue was washed with water, extracted with EtOAc, dried over anhydrous Na₂SO₄. The crude product was recrystallized from EtOAc.

5.1.5.1. Ethyl 4-(phthalazin-1-yl)piperazine-1-carboxylate (8a)

Yield (0.19 g; 68%), mp 271-273°C; ¹HNMR(300 MHz, DMSO): δ 8.10 (1H, s, phthalazine), δ 7.98-7.96 (2H, d, *J*= 7 Hz, phthalazine), δ 7.93-7.91 (2H, d, *J*= 7 Hz, phthalazine), δ 4.07 (2H, q,<u>CH₂-CH₃</u>), δ 3.63-3.62 (2H, m, piperazine ring), δ 2.86-2.85 (2H, m,piperazine ring), δ 2.55-2.54 (2H, m,piperazine ring), δ 2.27-2.26 (2H, m, piperazine ring), δ 1.14 (3H, t, CH₂-<u>CH₃</u>), FT-IR ($\dot{\nu}$ max, cm-¹): 1672 (C=O); MS (Mwt.: 286.33): m/z 286.05 (M⁺, 0.95%), 270.00 (3.77%), 56.00 (100.00%);Anal. Calcd for C₁₅H₁₈N₄O₂: C, 62.92; H, 6.34; N, 19.57; Found: C, 63.23; H, 6.38; N, 19.71

5.1.5.2. 1-(4-Phenylpiperazin-1-yl)phthalazine (8b)

Yield (0.20g; 69%), mp 242-244°C; ¹HNMR(300 MHz, DMSO): δ 8.15-8.13 (3H, m,phthalazine), δ 7.98-7.96 (2H, d, *J*= 7.5 Hz,phthalazine), δ 7.27-7.24 (1H, m,Ar-H), δ 6.98-6.95 (2H, m,Ar-H), δ 6.86-6.84 (2H, d, *J*= 7.2 Hz, Ar-H), δ 3.57-3.56 (2H, m,piperazine ring), δ 3.12-3.11 (2H, m,piperazine ring), FT-IR (iii max, cm⁻¹): 3428 (CH aromatic), 2991, 2919 (CH aliphatic); **MS** (Mwt.: 290.36): m/z 290.10 (M⁺, 0.41%), 162.10 (8.10%), 120.10 (100.00%),Anal. Calcd for C₁₈H₁₈N₄: C, 74.46; H, 6.25; N, 19.30; Found: C, 74.62; H, 6.37; N, 19.54.

5.1.5.3. 1-(4-(2-Fluorophenyl)piperazin-1-yl)phthalazine(8c)

Yield (0.21 g; 70%), mp 88-90°C; ¹HNMR(300 MHz, DMSO): δ 8.61 (1H, s,phthalazine), δ 8.27-8.24 (2H, d, J= 9 Hz, phthalazine), δ 8.14-8.11 (2H, d, J= 9 Hz, phthalazine), δ 7.19-7.17 (1H, d, J= 7.6 Hz, Ar-H), δ 7.15-7.13 (1H, d, J= 7.6 Hz, Ar-H), δ 7.08-7.06 (1H, d,J= 7.6 Hz, Ar-H), δ 6.99-6.97 (1H, d, J= 7.6 Hz, Ar-H), δ 3.70-3.69 (2H, m,piperazine ring), δ 3.31-3.30 (2H, m,piperazine ring), δ 3.23-3.21 (4H, m,piperazine ring); FT-IR (\acute{v} max, cm-¹): 3039 (CH aromatic), 2924 (CH aliphatic); MS (Mwt.: 308.35): m/z 308.05 (M⁺, 1.81%), 180.05 (19.88%), 138.10 (100.00%);Anal. Calcd for C₁₈H₁₇FN₄: C, 70.11; H, 5.56; N, 18.17; Found: C, 70.29; H, 5.60; N, 18.28.

5.1.5.4. 1-(4-(Pyridin-2-yl)piperazin-1-yl)phthalazine (8d)

Yield (0.20g; 69%), mp 286-288°C; ¹HNMR(300 MHz, DMSO): δ 8.14-8.12 (1H, d, *J*= 7 Hz,pyridine ring), δ 7.93-7.91 (3H, m,phthalazine), δ 7.59-7.56 (2H, d, *J*= 9 Hz, phthalazine), δ 7.56-7.54 (1H, d, *J*= 7 Hz,pyridine ring), δ 6.69-6.67 (2H, d, *J*= 7 Hz,pyridine ring), δ 3.67-3.65 (4H, m,piperazine ring), δ 3.06-3.04 (4H, m,piperazine ring); FT-IR (δ max, cm⁻¹): 3030 (CH aromatic), 2926 (CH aliphatic); **MS** (Mwt.: 291.35): m/z 291.10(M+, 0.19%), 279.05 (0.28%), 160.10 (27.24%), 95.05 (100.00); Anal. Calcd for C₁₇H₁₇N₅: C, 70.08; H, 5.88; N, 24.04; Found: C, 70.32; H, 5.64; N, 24.31

5.1.5.5. furan-2-yl(4-(phthalazin-1-yl)piperazin-1-yl)methanone (8e)

Yield (0.23g; 77%), mp 90-92°C; ¹HNMR(300 MHz, DMSO): δ 8.36 (1H,s, phthalazine), δ 8.12-8.10 (1H, d,*J*= 7 Hz, furoyl ring), δ 8.03-8.01 (2H, d, *J*= 7.8 Hz, phthalazine), δ 7.98-7.96 (2H, d, *J*= 7.8 Hz, phthalazine), δ 6.99-6.98 (2H, d, *J*= 7 Hz, furoylring), δ 3.69-3.67 (4H, m,piperazine ring), δ 2.91-2.90 (4H, m, piperazine ring); FT-IR (δ max, cm⁻¹): 1679 (C=O); **MS** (Mwt.: 308.33): m/z 308.05 (M+, 4.64%), 274.00 (6.66%), 158.05 (57.70%), 95.00 (100.00%) Anal. Calcd for C₁₇H₁₆N₄O₂: C, 66.22; H, 5.23; N, 18.17; Found: C, 66.41; H, 5.32; N, 18.40

5.1.5.6. Ethyl 4-(4-methylphthalazin-1-yl)piperazine-1-carboxylate (8f)

Yield (0.22 g; 73%), mp 95-97°C; ¹HNMR(300 MHz, CDCl₃): δ 8.43-8.41 (2H, d, *J*= 8.1 Hz, phthalazine), δ 8.36-8.34 (2H, d, *J*= 8.1 Hz, phthalazine), δ 4.19 (2H, q,<u>CH₂-CH₃), δ 3.75-3.74 (2H, m,piperazine ring), δ 3.65-3.64 (2H, m,piperazine ring), δ 3.37-3.36 (2H, m,piperazine ring), δ 3.18-3.17 (2H, m,piperazine ring), δ 2.58 (3H, s, CH₃), δ 1.33 (3H, t,<u>CH₃-CH₂</u>); FT-IR (δ max, cm⁻¹): 1693 (C=O); **MS** (Mwt.: 300.36): m/z 300.05(M+, 5.71%), 244.05 (3.15%), 184.55 (13.85%), 172.30 (48.47%), 55.65 (100%); Anal. Calcd for C₁₆H₂₀N₄O₂: C, 63.98; H, 6.71; N, 18.65; Found: C, 64.17; H, 6.83; N, 19.01</u>

5.1.5.7. 1-Methyl-4-(4-phenylpiperazin-1-yl)phthalazine (8g)

Yield (0.15g; 56%), mp 229-231°C; ¹HNMR(300 MHz, DMSO): δ 7.28-7.26(2H, d, *J*= 7.2 Hz, phthalazine), δ 7.25-7.23 (2H, d, *J*= 7.2 Hz, phthalazine), δ 6.99-6.97 (2H, d, *J*= 7.8 Hz, Ar-H), δ 6.88-6.86 (1H, m, Ar-H), δ 6.85-6.83 (2H, d, *J*= 7.8 Hz, Ar-H), δ 3.34-3.32 (4H, m, piperazine ring), δ 3.22-3.20 (4H, m, piperazine ring), δ 2.48 (3H, s,CH₃); FT-IR ($\dot{\nu}$ max, cm⁻¹): 3070 (CH aromatic), 2954 (CH aliphatic); MS (Mwt.: 304.39): m/z 304.00(M+, 1.73%), 284.60 (2.19%), 192.35 (5.24%), 105.20 (34.33%), 55.65 (100.00%); Anal. Calcd for C₁₉H₂₀N₄: C, 74.97; H, 6.62; N, 18.41; Found: C, 80.32; H, 6.70; N, 18.48.

5.1.5.8. 1-(4-(2-fluorophenyl)piperazin-1-yl)-4-methylphthalazine (8h)

Yield (0.12 g; 71%), mp 149-151°C; ¹HNMR(300 MHz, CDCl₃): δ 8.15-8.13 (1H, d, *J*= 9.6 Hz, phthalazine), δ 8.05-8.01 (1H, d, *J*= 9.6 Hz, phthalazine), δ 7.87-7.83 (2H, d, *J*= 9.6 Hz, phthalazine), δ 7.07-7.04 (2H, d, *J*= 8.7 Hz, Ar-H), δ 6.99-6.97 (2H, d,Ar-H), δ 3.69-3.67 (4H, m,piperazine ring), δ 3.48-3.46 (4H, m, piperazine ring), δ 2.03 (3H, s, CH₃); FT-IR ($\dot{\nu}$ max, cm⁻¹): 3035 (CH aromatic), 2951 (CH aliphatic); MS (Mwt.: 322.38): m/z 323.50 [M+1H]⁺ (9.74%), 172.15 (48.77%,), 122.05 (100.00%); Anal. Calcd for C₁₉H₁₉FN₄: C, 70.79; H, 5.94; N, 17.38; Found: C, 71.03; H, 5.99; N, 18.48.

5.1.5.9. 1-Methyl-4-(4-(pyridin-2-yl)piperazin-1-yl)phthalazine (8i)

Yield (0.10g; 59%), mp 226-228°C; ¹HNMR(300 MHz, DMSO): δ 8.16-8.13(1H, d, *J*= 8.7 Hz,pyridyl ring), δ 7.62-7.60 (2H, d, *J*= 8.4 Hz, phthalazine), δ 7.58-7.56 (2H, d, *J*= 8.4 Hz, phthalazine), δ 6.91-6.89 (1H, d, *J*= 8.7 Hz, pyridyl ring), δ 6.74-6.72 (2H, d, *J*= 8.7 Hz, pyridyl ring), δ 3.75-3.73 (4H, m, piperazine ring), δ 3.12-3.10 (4H, m,piperazine ring), δ 2.08 (3H, s,CH₃); FT-IR (δ max, cm⁻¹) 3132 (CH aromatic), 2954 (CH aliphatic); **MS** (Mwt.: 305.38): m/z 305.15 (M+, 1.16%), 185.65 (16.50%), 172.35 (41.23%), 55.55 (100.00%); Anal. Calcd for C₁₈H₁₉N₅: C, 70.80; H, 6.27; N, 22.93; Found: C, 71.04; H, 6.38; N, 23.18.

5.1.5.10. Furan-2-yl(4-(4-methylphthalazin-1-yl)piperazin-1-yl)methanone (8j)

Yield (0.11g; 60%), mp 90-92°C; ¹HNMR(300 MHz, DMSO): δ 8.32-8.30 (2H, d, *J*= 7 Hz, phthalazine), δ 8.29-8.27 (2H, d, *J*= 7 Hz, phthalazine), δ 7.82-7.81 (1H, d, *J*= 7.2 Hz, furoyl ring), δ 6.97-6.96(1H, d, *J*= 7.2 Hz, furoyl ring), δ 6.62-6.60 (1H, t, *J*= 7.2 Hz, furoyl- H), δ 3.64-3.62 (4H, m,piperazine ring), δ 3.41-3.39 (4H, m,piperazine ring), δ 2.48 (3H, s,CH₃); FT-IR (δ max, cm-¹): 1610 (C=O); **MS** (Mwt.: 322.36): m/z 322.00 (M+, 6.60%), 184.00 (31.70%), 172.00 (100.00%); Anal. Calcd for C₁₈H₁₈N₄O₂: C, 67.07; H, 5.63; N, 17.38; Found: C, 67.21; H, 5.69; N, 17.52

5.2. Biological Evaluation

5.2.1. Evaluation for cytotoxic activity against NCI 60 human cancer cell lines panel.

5.2.1.1 Assay protocol

The human tumor cell lines of the cancer-screening panel were grown in RPMI 1640 medium containing 5% fetal bovine serum and 2 μ M L-glutamine. For a typical screening experiment, cells are inoculated into 96 well microtiter plates in 100 ml at plating densities ranging from 5000 to 40,000 cells/well depending on the doubling time of individual cell lines. After cell inoculation, the microtiter plates are incubated at 37 °C, 5% CO₂, 95% air and 100%

relative humidity for 24 h prior to addition of experimental drugs. After 24 h, two plates of each cell line are fixed in situ with TCA, to represent a measurement of the cell population for each cell line at the time of drug addition (Tz). Experimental drugs are solubilized in dimethyl sulfoxide at 400-fold the desired final maximum test concentration and stored frozen prior to use. At the time of drug addition, an aliquot of frozen concentrate is thawed and diluted to twice the desired final maximum test concentration with complete medium containing 50 mg/ml Gentamicin. Additional four, 10-fold or ¹/₂log serial dilutions are made to provide a total of five drug concentrations plus control. Aliquots of 100 ml of these different drug dilutions are added to the appropriate microtiter wells already containing 100 ml of medium, resulting in the required final drug concentrations. Following drug addition, the plates are incubated for an additional 48 h at 37 °C, 5% CO₂, 95% air, and 100% relative humidity. For adherent cells, the assay is terminated by the addition of cold TCA. Cells are fixed in situ by the gentle addition of 50 ml of cold 50% (w/v) TCA (final concentration, 10% TCA) and incubated for 60 min at 4 °C. The supernatant is discarded, and the plates are washed five times with tap water and air dried. Sulforhodamine B (SRB) solution (100 ml) at 0.4% (w/v) in 1% acetic acid is added to each well, and plates are incubated for 10 min at room temperature. After staining, unbound dye is removed by washing five times with 1% acetic acid and the plates are air dried. Bound stain is subsequently solubilized with 10 µMtrizma base, and the absorbance is read on an automated plate reader at a wavelength of 515 nm. For suspension cells, the methodology is the same except that the assay is terminated by fixing settled cells at the bottom of the wells by gently adding 50 ml of 80% TCA (final concentration, 16% TCA).

5.2.1.2. Data analysis

Using the seven absorbance measurements [time zero, (Tz), control growth, (C), and test growth in the presence of drug at the five concentration levels (Ti)], the percentage growth is calculated at each of the drug concentrations levels. Percentage growth inhibition is calculated as: $[(Ti - Tz)/(C - Tz)] \times 100$ for concentrations for which Ti>/ ¹/₄ Tz and [(Ti -Tz) / Tz] x100 for concentrations for which Ti <Tz.

Three dose response parameters are calculated for each experimental agent. Growth inhibition of 50% (GI₅₀) is calculated from [(Ti -Tz)/(C - Tz)] x 100 =50, which is the drug concentration resulting in a 50% reduction in the net protein increase (as measured by SRB staining) in control cells during the drug incubation. The drug concentration resulting in total growth inhibition (TGI) is calculated from Ti ¹/₄ Tz. The LC₅₀ (concentration of drug resulting in a 50% reduction in the measured protein at the end of the drug treatment as compared to that at the beginning) indicating a net loss of

cells following treatment is calculated from [(Ti - Tz) / Tz] x100 = -50. Values are calculated for each of these three parameters if the level of activity is reached; however, if the effect is not reached or is exceeded, the value for that parameter is expressed as greater or less than the maximum or minimum concentration tested (M.C. Alley, 1988, M.R. Grever, 1992).

5.2.2. In vitro VEGFR tyrosine kinase activity at single dose of 10 µM concentration.

The VEGFR tyrosine kinase activity at single dose concentration of 10 μ M was carried out by BPS Bioscience (www.bpsbioscience.com). VEGFR (BPS#40301) served as the enzyme source and Poly (Glu, Tyr) sodium salt, (4:1, Glu:Tyr) (Sigma#P7244) served as the standardized substrate & Kinase-Glo Plus Luminescence kinase assay kit (Promega#V3772). The IC₅₀ determination was carried out where quality control testing is routinely performed on each of the targets to insure compliance to acceptable standards. ³³P-ATP was purchased from Perkin Elmer and ADP-GloTM was purchased from Promega. All other materials were of standard laboratory grade.

5.2.2.1. Assay protocols.

The VEGFR tyrosine kinase activity was performed using Kinase-Glo Plus luminescence kinase assay kit (Promega). It measures kinase activity by quantitating the amount of ATP remaining in solution following a kinase reaction. The luminescent signal from the assay is correlated with the amount of ATP present and is inversely correlated with the amount of kinase activity. The compounds were diluted to 100 µM in 10% DMSO and 5 ml of the dilution was added to a 50 ml reaction so that the final concentration of DMSO is 1% in all of reactions. All of the enzymatic reactions were conducted at 30 °C for 40 min. The 50 µl reaction mixture contains 40 µMTris, pH 7.4, 10 µM MgCl₂, 0.1 mg/ml BSA, 0.2 mg/ml Poly (Glu, Tyr) substrate, 10 µM ATP and VEGFR. After the enzymatic reaction, 50 ml of Kinase-GloPlus Luminescence kinase assay solution (Promega) was added to each reaction and incubate the plate for 5 min at room temperature. Luminescence signal was measured using a BioTek Synergy 2 microplate reader. The protein kinase assays used to determine IC₅₀ value were performed using ADP-GloTM assay kit from Promega which measures the generation of ADP by the protein kinase. Generation of ADP by the protein kinase reaction leads to an increase in luminescence signal in the presence of ADP-GloTM assay kit. The assay was started by incubating the reaction mixture in a 96-well plate at 30 °C for 30 min. After the 30 min incubation period, the assay was terminated by the addition of 25 ml of ADP-GloTM Reagent (Promega). The 96 well plate was shaken and then incubated for 40 min at ambient temperature. 50 ml of Kinase detection reagent was added, the 96well reaction plate was then read using the ADP-Glo Luminescences Protocol on a GloMax plate reader (Promega: Catalog #E7031). Blank control was set up that included all the assay components except the addition of appropriate substrate (replace with equal volume of kinase assay buffer). The corrected activity for each protein kinase target was determined by removing the blank control value.

5.2.2.2. Data analysis.

VEGFR activity assays were performed in duplicate at each concentration. The luminescence data were analyzed using the computer software, Graphpad Prism. The difference between luminescence intensities in the absence of VEGFR (Lu_t) and in the presence of VEGFR (Lu_c) was defined as 100% activity (Lu_t - Lu_c). Using luminescence

signal (Lu) in the presence of the compound, % activity was calculated as: % Activity = $\{(Lu_t - Lu)/(Lu_t - Lu_c) X100\%, where Lu = the luminescence intensity in the presence of the compound (all percent activities below zero were set to 0%). % Inhibition was calculated as: % inhibition =100 (%) - % activity. IC₅₀ determination for inhibitor against VEGFR was estimated by generating a graph of log inhibitor vs normalized response with variable using the Prism software.$

5.3. Docking studies.

All molecular modeling calculation and docking studies were carried out using Accelry's discovery studio 2.5 (Accelry's Discovery Studio 2.5, 2010). The X-ray crystal structure of the kinase domain of VEGFR in complex with its inhibitor (PDB code4ASD) was recovered RSCB protein data bank. Docking engine was validated; the co-crystallized ligand was extracted from the catalytic site and redocked to calculate the root mean square difference (RMSD)between the top docking pose and original crystallographic geometry. The amino acid residues were ionized using role based technique and the missing residues were completed. The protein structure was typed by CHARMM force field then it was minimized using 500 step of SMART minimizer of Discovery studio program (Minimizer adopt hybrid of steepest descent and conjugate gradient minimization algorithms). All the water molecules in the protein were deleted. Then minimization was done to the ligand before docking to give the lowest conformational energy for the ligand after preparation step. Docking was performed using CDocker protocol with the kinase active site.

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"تصميم وتشييد مركبات مبنية على حلقة الفثلازين كمضادات محتملة للأورام السرطانية". للسادة الدكاتر ة

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يعتبر مرض السرطان هو السبب الرئيسي للوفاة في جميع أنحاء العالم. وكشفت الدراسات الوبائية أن السرطان يمثل واحدة من كل خمس حالات وفاة. وعلاوة على ذلك، تشير التقديرات إلى أن العدد السنوي للوفيات بسبب السرطان سيرتفع من ٧.٦ مليون في عام ٢٠٠٨ إلى ١٣ مليون في عام ٢٠٣٠. وعلى الرغم من تحسن التقنيات التشخيص الجزيئي، والتقدم في الوقاية والعلاج الكيميائي، ولكن يزال هذا المرض يؤثر على الملايين من المرضى في جميع أنحاء العالم. يعتبر نمو أوعية دموية جديدة هي واحدة من السمات المميزة الراسخة في عملية التسرطن. وكذلك يعتبر مستقبل (VEGFR) يلعب دورا حاسما في الأوعية الدموية السرطانية. من خلال استهداف مستقبل (VEGFR).

يحتوى هذا البحث على مقدمة تشمل نبذة عن مرض السرطان وعن الدور الفعال الذى يلعبه انزيم التيروزين كينيز بأنواعه المختلفة فى تطور المرض وكيفية تثبيط هذا الانزيم بواسطة مشتقات الأدوية المختلفة وكذلك استهداف مستقبل VEGFR-2 .

وفي هذا البحث، تم تصميم سلسلة جديدة من المركبات التى تحتوى على حلقة الفثلازين لإستهداف مستقبل VEGFR-2 كمثبطات وكذلك كمحفظ لموت الخلايا المبرمج. ولذلك تم تصميم ثلاثة مجموعات من مشتقات الفثلازين كمثبطات لإنزيم التيروزين كينيز وتم تقييم فاعليتها المصادة للأورام السرطانية. كما تم تدعيم هذا التصميم بدراسة النمذجة الجزيئية بإستخدام برنامج "C-Docker" عن طريق عمل إرساء للمركبات المقترحة على مستقبل إنزيم التيروزين كينيز(2-70)

وقد تم إثبات صحة التركيب الكيميائى لهذه المركبات عن طريق الأشعة تحت الحمراء والرنين النووى المغناطيسي لنواة الهيدروجين (١) وطيف الكتلة والتحليل الدقيق للعناصر المكونة لها.