Supplementary Information

Discovery of new 1,3-diphenylurea appended aryl pyridine derivatives as apoptosis inducers through c-MET and VEGFR-2 inhibition: Design, synthesis, *in vivo* and *in silico* studies

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Figure S1. ¹H NMR (500 MHz, DMSO-*d*₆) spectrum of compound 2a





Figure S3. Mass spectrum of compound 2a



Figure S4. IR spectrum of compound 2a







Figure S6. ¹³C NMR (125 MHz, DMSO-*d*₆) spectrum of compound 2b



Figure S7. Mass spectrum of compound 2b







Figure S9. ¹³C NMR (125 MHz, DMSO- d_6) spectrum of compound 2c



Figure S10. Mass spectrum of compound 2c



Figure S11. ¹H NMR (500 MHz, DMSO-*d*₆) spectrum of compound 2d



Figure S12. ¹³C NMR (125 MHz, DMSO-*d*₆) spectrum of compound 2d







Figure S14. IR spectrum of compound 2d



Figure S15. ¹H NMR (500 MHz, DMSO-*d*₆) spectrum of compound 2e



Figure S16. ¹³C NMR (125 MHz, DMSO- d_6) spectrum of compound 2e



Figure S17. Mass spectrum of compound 2e



Figure S18. IR spectrum of compound 2e



Figure S19. ¹H NMR (500 MHz, DMSO-*d*₆) spectrum of compound 2f



Figure S20. ¹³C NMR (125 MHz, DMSO-*d*₆) spectrum of compound 2f



Figure S21. Mass spectrum of compound 2f



Figure S22. ¹H NMR (500 MHz, DMSO-*d*₆) spectrum of compound 2g



Figure S23. ¹³C NMR (125 MHz, DMSO-*d*₆) spectrum of compound 2g



Figure S24. Mass spectrum of compound 2g



Figure S25. ¹H NMR (500 MHz, DMSO-*d*₆) spectrum of compound 2h



Figure S26. ¹³C NMR (125 MHz, DMSO-*d*₆) spectrum of compound 2h



Figure S27. Mass spectrum of compound 2h



Figure S28. ¹H NMR (500 MHz, DMSO-*d*₆) spectrum of compound 2i



Figure S29. ¹³C NMR (125 MHz, DMSO- d_6) spectrum of compound 2i



Figure S30. Mass spectrum of compound 2i



Figure S31. ¹H NMR (500 MHz, DMSO-*d*₆) spectrum of compound 2j



Figure S32. ¹³C NMR (125 MHz, DMSO-*d*₆) spectrum of compound 2j



Figure S33. Mass spectrum of compound 2j



Figure S34. ¹H NMR (500 MHz, DMSO-*d*₆) spectrum of compound 2k



Figure S35. ¹³C NMR (125 MHz, DMSO-*d*₆) spectrum of compound 2k



Figure S36. Mass spectrum of compound 2k



Figure S37. ¹H NMR (500 MHz, DMSO-*d*₆) spectrum of compound 2l



Figure S38. ¹³C NMR (125 MHz, DMSO-*d*₆) spectrum of compound 2l



Figure S39. Mass spectrum of compound 21



Figure S40. IR spectrum of compound 21



Figure S41. ¹H NMR (500 MHz, DMSO-*d*₆) spectrum of compound 2m



Figure S42. ¹³C NMR (125 MHz, DMSO-*d*₆) spectrum of compound 2m



Figure S43. Mass spectrum of compound 2m



Figure S44. ¹H NMR (500 MHz, DMSO-*d*₆) spectrum of compound 2n



Figure S45. ¹³C NMR (125 MHz, DMSO-*d*₆) spectrum of compound 2n



Figure S46. Mass spectrum of compound 2n

RequesterData						
Name:	Haytham O. Tawfik	Tel.	01005356819			
Authority:	Faculty of Pharmacy Tanta University	Date	30/03/2023			

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 $Samples had been {\it submitted for elemental analysis}.$

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No.	Code	C%	H%	N%
1	2a	73.60	4.50	13.66
2	2b	68.38	4.01	12.59
3	2 c	62.11	3.44	11.44
4	2d	73.95	4.91	13.45
5	2e	71.33	4.50	12.77
6	2f	66.69	3.75	15.64
7	2g	63.01	3.49	11.87
8	2h	63.00	3.33	11.70
9	2i	62.98	3.45	11.80
10	2j	52.01	2.91	9.54
11	2k	67.99	4.99	11.35
12	21	76.54	4.52	12.40
13	2m	70.98	4.25	17.25
14	2n	72.52	4.18	15.85
INVESTIGATOR DIRECTOR				

Figure S47. Elemental analysis of compounds 2a-n

HPLC Protocol Data

The lead compounds were tested for purity using HPLC (AgilentTM 1260 Infinity HPLC system equipped with Agilent 1260 Infinity quaternary pump (G1311C), Agilent 1260 Infinity thermostated column compartment (G13b16A), Agilent 1260 Infinity autosampler with reliable injections from 0.1 to 100 μ L (G1329B) and Agilent 1260 Infinity UV detector (G1321C). Data was recorded and analyzed with Agilent OpenLAB CDS ChemStation Edition software. HANNA pH 211 Microprocessor pH meter with double junction glass electrode was used to adjust the pH. The column used in HPLC was ZORBAX Eclipse plus C18 (250 x 4.6 mm, 3.5 μ m). Mobile phase consisted of Acetonitrile (ACN)/0.05M KH2PO4 buffer pH3.7 (30:70), flow rate of 1.5 mL/min and the run time was set as double the elution time of each compound. Their purity was recorded with a high percentage (\geq 95).

Acq. Operator	:	SYSTEM
Sample Operator	:	SYSTEM
Acq. Instrument	:	HPLC Location : Vial15
Injection Date	:	8/23/2023 1:28:39 PM
		Inj Volume : 5.000 μl
Method	:	C:\CHEM32\1\METHODS\DEF_LC.MCYANO COLUMN.M
Last changed	:	8/23/2023 2:39:15 PM by SYSTEM
		(modified after loading)
Sample Info	:	30 ACN:70 phosphate buffer pH 3.7, Flow 1.50 mL/min, 254 nm, 5 ul injection



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Area Percent Report
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Sorted By Calib. Data Modified Multiplier	:::::::::::::::::::::::::::::::::::::::	Signal 8/23/2023 2:39:1 1.0000	5 PM	
Sample Amount:	•	1.0000	[(not used in calc.)
Do not use Multiplier	&	Dilution Factor with	[mg/mi] ISTDs	(not used in carc.)

Signal 1: VWD1 A, Wavelength=254 nm

Totals : 2.06465e4 98.5384

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*** End of Report ***
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Figure S48. HPLC chart of compound 2n



Figure S49. Inhibition ratio curves of 2a-n towards c-Met. Different compound concentrations were set in the assay.



Figure S50. Inhibition ratio curves of 2a-n towards VEGFR-2. Different compound concentrations were set in the assay.

Kinase	Kinase activity (%)	Kinase inhibitory activity (%)
p38a MAPK	98	2
SGK1	93	7
GSK3b	104	-4
Aurora B	103	-3
MARK3	85	15
Src	89	11
BTK	97	3

Table S1. Percent kinase activity and inhibitory values of compound 2n against a panel of kinases

Biology Protocols

c-MET and VEGFR-2 kinase inhibition Assays

All molecules were evaluated for their c-MET (BPS Bioscience Corporation catalog #79559) and VEGFR-2 kinase inhibition using (BPS Bioscience Corporation catalog#40325) using ELISA kit (Enzyme-Linked Immunosorbent Assay). They were solvated in DMSO (0.1%), and serial diluations (four conc.) were set following the manufacturer's instructions.

Cytotoxicity

Both breast cancer "MCF-7" and prostate cancer "PC-3" and normal "WISH" cells were purchased from the National Research Institute, Egypt, and maintained in RPMI-1640 medium L-Glutamine (Lonza Verviers SPRL, Belgium, cat#12-604F). All cells were incubated at 37 °C in a 5% carbon dioxide atmosphere (NuAire). Cells were plated at a density of 5×10^4 cells in triplicates in a plate of 96 wells. On the second day, cells were treated with the compounds with concentrations of (6.25, 12, 25, 50, and 100 µM). Cell viability was assessed using the MTT assay.

Investigation of apoptosis

Annexin V/PI staining and cell cycle analysis

MCF-7 cells were seeded into 6-well culture plates $(3-5 \times 10^5 \text{ cells/well})$ and incubated overnight. Cells were treated with compound **APPU 2n** at their IC₅₀ values for 48 h. Next, media supernatants and cells were collected and rinsed with ice-cold PBS. Then, cells were suspended the cells in 100 µL of annexin binding buffer solution "25 mM CaCl₂, 1.4 M NaCl, and 0.1 M Hepes/NaOH, pH 7.4" and incubated with "Annexin V-FITC solution (1:100) and propidium iodide (PI)" at a concentration equals 10 µg/mL in the dark for 30 min. Stained cells were then acquired by BD FACSCaliburTM Flow Cytometer.

Real-time-polymerase chain reaction for the selected genes

Gene expression of Bcl-2, the anti-apoptotic gene, and the pro-apoptotic genes "P53, Bax, and Caspapses-3,8, and 9" were examined to delve deeper into the apoptotic pathway. MCF-7 cells were then treated with compound **APPU 2n** at their IC₅₀ values for 48 h. After treatment, the RT-PCR reaction was carried out following routine

work. Then, the Ct values were collected to calculate the relative genes' expression in all samples by normalization to the β -actin housekeeping gene.

In Vivo Assay

Animals and tumor cell line

Adult female Swiss albino mice purchased from Theodor Bilharzia Research Institute, Giza, Egypt, with an average body weight of (18-23) g was used. Mice were housed under constant conditions of 12 h light/dark cycle in a temperature under conditions of controlled humidity (22 ± 2 °C), with free access to standard laboratory mice food and water.

Solid Ehrlich carcinoma (SEC) cells were got from the National Cancer Institute (Cairo University, Egypt). The tumor cell line was proliferated in mice through serial intraperitoneal (I.P.) transplantation of a volume of 0.2 mL physiological saline containing 1×10^6 viable cells for 24 h. SEC cells were collected 7 days after I.P. implantation. The harvested cells were diluted with saline to obtain a concentration of 5×10^6 viable SEC cells/mL. A volume of 0.2 mL saline contains 1×10^6 SEC cells that were I.P. implanted into each normal mouse. SEC cells (1×10^6 tumor cells/mouse) were implanted subcutaneously into the right thigh of the hind limb.

The experimental animals were randomly divided into four groups. Group 1 served as the normal saline control. Group 2 served as the SEC control (1×10^6 cells/mouse). Group 3 served as the compound **2n**-treated group (6 mg/kg B.Wt., I.P.). Group 4 received the standard anticancer drug of Cabozantinib (6 mg/kg BW, I.P.) and is considered as a reference control. Body weight and survival were recorded daily until the 24th day in both treated and control groups. At the end of experiment, the blood of each group was collected under light anesthesia for estimation of hematological assays. The anesthetized animals were then sacrificed for evaluation of the antitumor activity examination.

Antitumor potentiality

It includes tumor volume, weight, and tumor inhibition ratio (TIR%). Time interval measurements of tumor volume using digital Vernier caliper (Tricle Brand, Shanghai, China). Measure tumor length and width using clipper and then calculate tumor volume using formulations $V = (L \times W \times W)/2$, where V is tumor volume, W is tumor width, L is tumor length. While TIR% was calculated according to the following equation $\frac{Tumor volume (Control) - Tumor volume (treated)}{T} \times 100$.

Tumor volume (control)

Blood assays

At the end of the experiment, animals from different groups were sacrificed, and blood samples were collected for determination of was estimated using Abbott CELL-DYN® 1800 automated hematology analyzer (USA) with ready-made kits (Abbott Laboratories, Abbott Park, IL, USA).



Figure S51. Overlay of the docked ligand (green) with the co-crystalized ligand (red) inside the active site of A) c-MET (RMSD= 1.0906 Å) and B) VEGFR-2 (RMSD= 1.0042 Å) receptors.



Figure S52. From left to right: (A) Temperature, (B) pressure and (C) potential energy during the 100ns MD simulations (Top: VEGFR-2 system and Bottom: c-MET system).



Figure S53. Frequency of interacting residues and type of interaction for 2n with VEGFR-2 and c-MET receptors.