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Synthesis, Anticancer and Molecular Docking Studies of 2-(4-chlorophenyl)-5-aryl-1,3,4-Oxadiazole Analogues

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Introduction

About 13 percent of all the death worldwide is due to cancer, surpassing cardiovascular disease and taking number one place [1,2]. Chemotherapy of cancer is associated with various adverse effects viz. bone marrow depression, alopecia, drug induced caner, etc. and is often associated with cytotoxicity, genotoxicity to normal cells together with the development of resistance [3]. Medicinal chemists have great perseverance in research and development (R & D) for the search of newer and safer anticancer agents. EGFR family of Tyrosine Kinases (TK) play a vital role in cancer proliferation and it is suggested that any agent which would inhibit the TK activity may have substantial role in the cancer treatment [4]. So we selected EGFR family of TK and explore the binding mode of the our compounds to EGFR tyrosine kinase active site. Imatinib (gleevec) an anticancer drug is TK inhibitor (TKI), inhibits TK encoded by the bcr-abl oncogene as well as receptor TKs encoded by the c-kit and platelet-derived growth factor receptor (PDGFR) oncogenes [5]. Oxadiazole derived compounds are known to display wide range of biological and pharmacological activities including anticancer, antitubercular, antibacterial, antifungal, anti-HIV, anti-inflammatory, and insecticidal activities [6-12]. There are nearly 2577 publications from 2002 to 2012 involving 1,3,4-oxadiazoles [13]. Some of the marketed oxadiazole drugs include raltegravir (antiretroviral), zibotentan (anticancer), etc. We have earlier reported the anticancer activity of some novel oxadiazole analogues [6,14].

Materials and Methods

Chemistry

All chemicals were supplied by E. Merck, and S. D. Fine Chemicals.

Melting points were determined by open tube capillary method and are uncorrected. Purity of the compounds was checked by elemental analysis and the progress of reactions was monitored by TLC plates (silica gel G) using mobile phase, chloroform: methanol (9:1), and acetone: n-hexane (8:2) and the spots were identified by iodine vapours or UV light. IR spectra were recorded on a Schimadzu 8201 PC, FT-IR spectrometer (KBr pellets). 1H NMR spectra were recorded on a Bruker AC 300 MHz spectrometer using TMS as internal standard in DMSO *d6*. Mass spectra were recorded on a Bruker Esquire LCMS using ESI and elemental analyses were performed on Perkin-Elmer 2400 Elemental Analyzer.

General method for the synthesis of 4-chlorobenzohydrazide (3): 4-Chlorobenzoic acid (1) (7.84 g, 0.05 mol) was dissolved in excess of ethanol (50 ml) the reaction mixture was acidified and refluxed for 8-10 h. The layer of ester is separated by filtration flask and neutralized with sodium bicarbonate to obtain ethyl-4-chlorobenzoate (2). Equimolar mixture of ethyl-4-chlorobenzoate (2) and hydrazine hydrate was

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refluxed for 12 h and the excess solvent removed under vacuum and poured into the crushed ice to obtain 4-chlorobenzohydrazide (3).

General method for the synthesis of 2-(4-chlorophenyl)-5-aryl-1,3,4-oxadiazole analogues(4a-j): 4-Chlorobenzohydrazide (0.85 g, 0.005 mol) (3) and aromatic aldehydes was refluxed 10-12 h using 20 mol% NaHSO3 and ethanol-water system (1:2, v/v) solvent [15]. After completion of reaction the mixture the excess solvent removed and the concentrate was poured into crushed ice washed with water, dried and recrystallized with absolute ethanol. The reaction was monitored throughout by TLC using chloroform-methanol (9:1) and acetone: n-hexane (8:2) as mobile phase.

2-(4-Chlorophenyl)-5-(4-fluorophenyl)-1,3,4-oxadiazole (4a) Yield 70%, IR (KBr) 1521, 1112, 789, 745 cm-1; 1H NMR (300 MHz, DMSO-*d*6): δ 7.02-7.03 (2H, d, *J*=3.2 Hz, ArH), 7.36-7.38 (2H, d, *J*=6.1 Hz, ArH), 7.39-7.41 (2H, d, *J*=6.0 Hz, ArH), 7.43-7.45 (2H, d, *J*=6.2 Hz, ArH). 13C NMR (75 MHz, DMSO-*d*6): δ 116.1, 121.9, 124.5, 128.7, 129.2, 129.7, 134.7, 162.6, 164.8; *m*/*z*=274 (M+), 275 (M+1)+1, 276 (M+2)+. Cal/Ana: [C (61.12) 61.22 H (2.92) 2.94 N (10.08) 10.20].

2-(4-Chlorophenyl)-5-(4-chlorophenyl)-1,3,4-oxadiazole (4b) Yield 74%, IR (KBr) 1531, 1131, 742 cm-1; 1H NMR (300 MHz, DMSO-*d*6): δ 7.28-7.30 (4H, dd, *J*=6.1 Hz ArH), 7.39- 7.41 (4H, dd, *J*=6.2 Hz ArH); *m/z*=290 (M+), 292 (M+2)+. Cal/Ana: [C (57.62) 57.76 H (2.79) 2.77 N (9.59) 9.62].

2-(4-Chlorophenyl)-5-(4-methoxyphenyl)-1,3,4-oxadiazole (4c) Yield 82%, IR (KBr) 1526, 1121, 749 cm-1; 1H NMR (300 MHz, DMSO-*d*6): δ 3.79 (3H, s, OCH3), 7.32-7.34 (2H, d, *J*=6.1 Hz, ArH), 7.35-7.37 (2H, d, *J*=6.1 Hz, ArH), 7.39-7.41 (2H, d, *J*=6.0 Hz, ArH), 7.81- 7.83 (2H, d, *J*=6.1 Hz ArH); *m*/*z*=286 (M+), 288 (M+2)+. Cal/ Ana: [C (62.79) 62.84 H (3.89) 3.87 N (9.57) 9.77].

2-(4-Chlorophenyl)-5-(3,4-dimethoxyphenyl)-1,3,4-oxadiazole (4d) Yield 79%, IR (KBr) 1528, 1127, 694 cm-1; 1H NMR (300 MHz, DMSO-d6): δ 3.81 (6H, s, OCH3), 7.31-7.33 (2H, d, *J*=6.1 Hz, ArH), 7.42-7.44 (2H, d, *J*=6.2 Hz, ArH), 7.89-7.92 (3H, m, ArH); *m/z*=316 (M+), 318 (M+2)+. Cal/Ana: [C (60.47) 60.67 H (4.19) 4.14 N (8.77) 8.84].

2-(4-Chlorophenyl)-5-(4-hydroxy-3-methoxyphenyl)-1,3,4oxadiazole (4e) Yield 65%, IR (KBr) 3397, 1525, 1132, 699 cm-1; 1H NMR (300 MHz, DMSO-*d*6): δ 3.83 (3H, s, OCH3), 7.31-7.33 (2H, d, *J*=6.0 Hz, ArH), 7.42-7.44 (2H, d, *J*=6.1 Hz, ArH), 7.84-7.87 (3H, m, ArH), 10.27 (1H, s, OH); *m*/*z*=302 (M+), 304 (M+2)+. Cal/Ana: [C (59.47) 59.52 H (3.59) 3.66 N (9.22) 9.25].

2-(4-Chlorophenyl)-5-(4-hydroxyphenyl)-1,3,4-oxadiazole (4g) Yield 80%, IR (KBr) 3401, 1527, 1121, 699 cm-1; 1H NMR (300 MHz, DMSO-*d*6): δ 6.79-6.81 (2H, s, ArH), 7.27-7.29m (2H, d, *J*=6.0 Hz, ArH), 7.34-7.36 (2H, d, *J*=6.0 Hz, ArH), 7.41-7.43 (2H, d, *J*=6.1 Hz, ArH), 7.81-7.84 (3H, m, ArH), 10.12 (1H, s, OH); *m/z*=302 (M+), 304 (M+2)+. Cal/Ana: [C (59.45) 59.52 H (3.65) 3.66 N (9.28) 9.25].

2-(4-Chlorophenyl)-5-phenyl-1,3,4-oxadiazole (4h) Yield 66%, IR (KBr) 1529, 1137, 702 cm-1; 1H NMR (300 MHz, DMSO-*d*6): δ 7.31-7.33 (2H, d, *J*=6.0 Hz, ArH), 7.41-7.43 (2H, d, *J*=6.1 Hz, ArH), 7.67-7.71 (5H, m, ArH); *m*/*z*=256 (M+), 258 (M+2)+. Cal/Ana: [C (65.47) 65.51 H (3.59) 3.53 N (10.82) 10.91].

2-(4-Chlorophenyl)-5-(4-methylphenyl)-1,3,4-oxadiazole (4i) Yield 13%, IR (KBr) 1519, 1139, 714 cm-1; 1H NMR (300 MHz, DMSO-*d*6): δ 2.32 (3H, s, CH3), 7.14-7.16 (2H, d, *J*=6.0 Hz, ArH), 7.32-7.34 (2H, d, *J*=6.1 Hz, ArH), 7.35-7.37 (2H, d, *J*=6.0 Hz, ArH), 7.41- 7.43 (2H, d, *J*=6.0 Hz, ArH); *m*/*z*=270 (M+), 272 (M+2)+. Cal/ Ana: [C (66.47) 66.55 H (4.19) 4.10 N (10.31) 10.35].

2-(4-Chlorophenyl)-5-(furan-2-yl)-1,3,4-oxadiazole (4j) Yield 69%, IR (KBr) 1525, 1124, 731 cm-1; 1H NMR (300 MHz, DMSO-d6): δ 6.98-7.01 (3H, s, Furan), 7.32-7.34 (2H, d, *J*=6.0 Hz, ArH), 7.39-7.41 (2H, d, *J*=6.1 Hz, ArH); *m*/*z*=246 (M+), 248 (M+2)+. Cal/Ana: [C (58.37) 58.43 H (2.81) 2.86 N (11.32) 11.36].

Anticancer activity

The compounds (4a and 4c) submitted to the NCI 60 cell screen were tested initially at a single high dose (10-5 M) on leukemia, melanoma, lung, colon, CNS, ovarian, renal, prostate, and breast cancers cell lines, nearly 60 in number. The one-dose data was reported as a mean graph of the percent growth of treated cells. The number reported for the one-dose assay is growth relative to the no-drug control, and relative to the time zero number of cells. The anticancer screening was carried out as per the NCI US protocol reported elsewhere [16-19]. We have discussed the anticancer screening method in our previous work [6,14,20].

Molecular docking studies

X-ray crystal structure of EGFR tyrosine kinase (PDB: 2J5F) was downloaded from www.rcsb.org. The active site of 2J5F is well established with hydrophobic active site containing irreversible inhibitor and molecular docking simulations were performed in order to distinguish the basic receptor-ligand interactions. The X-ray crystal structure of EGFR tyrosine kinase domain had the resolution of 3.00Å. The protein was prepared by using the Protein Preparation Wizard, pre-processed and heterostate for co-crystallized ligand was generated using Epik; protonation state and optimization of H bonding of the protein side chains were assigned using Protassign, energy minimized (impref minimization) using OPLS2001 force field. Receptor grid has been prepared with default parameters and without any constrains. Site was specified around the reference ligands N-[4-(3- bromophenylamino)quinazolin-6-yl]acrylamide of EGFR tyrosine kinase. The three dimensional structures of ligands were drawn by using the Maestro 8.5. The ligands were prepared by using Ligprep utility of Schrodinger Suite with default parameters, the ligand energy minimized by using OPLS 2005 (Macromodel multiple minimization) and water as solvent. The ligands did not show the formation of any tautomers or isomers after ligprep and macromodel multiple energy minimizations. The ligands' docking was performed with Xtra precision mode (XP) which is employed in GLIDE 5.0 module implemented in the Schrodinger LLC.

Results and Discussion

Chemistry

In the first step 4-chlorobenzoic acid (1) in excess of ethanol was refluxed for 8-10 h in acidic medium to obtain ethyl-4-chlorobenzoate (2). In the subsequent step compound (2) was refluxed with hydrazine hydrate in ethanol for 12 h to obtain 4-chlorobenzohydrazide (3). In the final step 4-chlorobenzohydrazide (3) and aromatic aldehydes was refluxed 10-12 h using 20 mol% NaHSO3 and ethanol-water system (1:2, v/v) solvent to obtain oxadiazole analogues (4a-n). The reaction was monitored throughout by thin layer chromatography (TLC) using chloroform-methanol (9:1) and acetone: n-hexane (8:2) as mobile phase

and the purity of the compounds was checked by elemental analysis. The reaction sequence is shown in Scheme 1. The synthesized compounds were characterized by spectral analysis and all the compounds were in full harmony with the proposed structures. In general the IR spectra afforded absorption 1519-1531 cm-1 band due to C=N and 1112-1271 cm-1 due to oxadiazole stretching. In 1H NMR the signals of the respective protons of the synthesized title compounds were verified on the basis of their chemical shifts and multiplicities in DMSO *d6*. The spectra showed a singlet at δ 3.79-3.83 ppm corresponding to OCH3; a doublet or multiplet at δ 6.79-7.92 ppm corresponding to aromatic protons.

Anticancer activity

2-(4-chlorophenyl)-5-(4-fluorophenyl)-1,3,4-oxadiazole (4a) showed growth percent (GP) of 87.27 (SF-295; CNS cancer) and 89.12 (MCF7; Brest Cancer) while 2-(4- chlorophenyl)-5-(4-methoxyphenyl)-1,3,4-oxadiazole (4c) showed GP of 71.70 (PC-3; Prastate cancer) and 74.14 (SR; Leukemia). The compound 4c (mean GP; 95.37) was found to be more active than the compound 4a (mean GP; 98.74). The *in vitro* anticancer activity of the compounds is given in Table 1. The compound 4c with 4-methoxyphenyl at the 5 position of the oxadiazole ring showed more anticancer activity than the compound 4a with 4-fluorophenyl at the 5 position of oxadiazole nucleus.

Molecular docking studies

The EGFR tyrosine kinase was reported several times as target for the inhibition of cancer cells. It contains the bound ligand N-[4-(3-bromophenylamino) quinazolin-6-yl] acrylamide and is well established by the presence of hydrophobic cavity at active site. Redocking of bound or reference ligand 34-JAB with EGFR tyrosine kinase exhibited the hydrophobic interactions with the residues Thr790, Met793 and Cys797. The amino acid residues Lys745, Glu762, Met766, Leu788, Met793 and Thr854 make the receptor hydrophobic in nature. The most important residue Cys797 is present near to the para substitution of phenyl group. The five member oxadiazole ring of ligands was lying near to Leu792 and Met 793. It was observed that the presence of methoxy functional group at para position may become more selective towards the EGFR tyrosine kinase [14]. In order to predict the binding affinity and pre eminent docked structures, the combined ligand docking and energy-grid scores were ranked by using E model and Glide scores. The ligand docking and E model scores were provided in the Table 2. The docking scores of compounds 4a and 4c were -5.251 and -5.433 respectively. The molecular docking and binding of ligands 4a and 4c is shown in Graphical abstract, while the 2D pose of the ligand 4c and the active site EGFR tyrosine kinase is shown in Figures 1 and 2.

Conclusion

A series of 10 oxadiazole analogues were synthesized in satisfactory yield and two compounds were evaluated for their *in vitro* anticancer activity at single-dose assay. The oxadiazole analogues showed moderate anticancer activity on various cell lines and molecular docking studies showed that the residue Cys797 is present near to the para substitution of phenyl group while the five member oxadiazole ring of ligands was lying near to Leu792 and Met 793 of EGFR tyrosine kinase active site. The oxadiazole analogues reported in this study may be further modified to increase their anticancer activity.



Scheme 1: Protocol for the synthesis of 2-(4-chloro)-5-aryl-1,3,4-oxadiazole analogues (4a-j).

Comp.	60 cell lines assay in 1 dose 10⁵M conc.				
	NSC Code	Mean GP	Range of GP	The most sensitive cell line	GP of the most sensitive cell line
4a	776718	98.74	87.26 to 112.87	SF-295 (CNS Cancer)	87.26
				MCF7 (Brest Cancer)	89.12
				UO-31 (Renal Cancer)	89.93
				HCT-15 (Colon cancer)	90.55
				NCI-H522 (Non-Small Cell Lungs cancer)	91.27
4c	776717	95.37	71.70 to 110.76	PC-3 (Prostate Cancer)	71.70
				SR (Leukemia)	74.14
				UO-31 (Renal Cancer)	80.62
				NCI-H522 (Non-Small Cell Lungs cancer)	82.58
				SK-OV-3 (Ovarian Cancer)	83.34

Table 1: Anticancer activity of the selected oxadiazole analogues.



Figure 1: Superimposed diagram of the two ligands 4a and 4c (ball and stick model).



Figure 2: (A) ligand 4c was shown in ball and stick model and remaining were EGFR tyrosine kinase active site residues; (B) 2D pose view diagram of the ligand 4c and EGFR tyrosine kinase active site residues.

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S. No.	Compound	Docking score	E-model score
1	Reference [21]	-8.288	-68.491
2	4a	-5.251	-37.778
3	4c	-5.433	-38.804

 Table 2: The docking score and E model score of reference ligand and selected ligands (4a and 4c).

The authors confirm that this article content has no conflicts of interest.

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