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Synthesis and molecular structure of arene ruthenium(II) benzhydrazone complexes: Impact of substitution at chelating ligand and arene moiety on antiproliferative activity

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Abstract
Convenient method of synthesis of ruthenium(II) arene benzhydrazone complexes (1-6) of the general formula [(η⁶-arene)Ru(L)Cl] (arene-benzene or p-cymene; L-monobasic bidentate substituted indole-3-carboxaldehyde benzhydrazone derivatives) has been described. The complexes have been fully characterized by elemental analysis, IR, UV-vis, NMR and ESI-MS spectral methods. The solid state molecular structures of representative complexes were determined by a single-crystal X-ray diffraction study and it indicates the presence of pseudo octahedral (piano stool) geometry. All the complexes were thoroughly screened for their cytotoxicity against human cervical cancer cells (HeLa), human breast cancer cell line (MDA-MB-231) and human liver carcinoma cells (Hep G2) under in vitro conditions. Interestingly, the cytotoxic activity of the complexes 3, 4 and 6 are much more potent than cis-platin with low IC₅₀ values against all the cancer cell lines tested. Further, the mode of cell death in MDA-MB-231 cells was assessed by AO-EB staining, Hoechst 33258 staining and flow cytometry technique along with comet assay. Further, the result of Western blot analysis suggests that complexes 3 and 6 were shown to accumulate preferentially in the mitochondria of MDA-MB-231 cells and induce apoptosis via mitochondrial pathway by up-regulating of p53 and Bax, and down-regulating Bcl-2.

Introduction
Over the past few decades, a large number of cisplatin analogs have been screened as potential antitumor agents, but of these only two, carboplatin and oxaliplatin, have entered world-wide clinical use.¹ Regardless of the achievements of current platinum drugs, they are efficient only for a limited range of cancers or some tumors can have acquired or intrinsic resistance, and they often cause severe side-effects.²,³ Hence, there is a need for new
approaches that are purposefully designed to circumvent these drawbacks. In this regard, ruthenium compounds in +2 or +3 oxidation state are considered to be suitable candidates for anticancer drug design, since they exhibit a similar spectrum of kinetics for their ligand substitution reactions as platinum(II). A number of ruthenium compounds have been shown to display promising anticancer activity and two ruthenium(III) complexes have entered clinical trials, trans-[RuCl₄(DMSO)(Im)]ImH (NAMI-A, where Im-imidazole),⁴ and trans-[RuCl₄(Ind)₂]IndH (KP1019), where Ind-indazole.⁵

Several reports have been focused on the anticancer potential of half-sandwich Ru(II) arene complexes of the type, [(η⁶-arene)Ru(YZ)(X)], where Y and Z are bidentate chelating groups (NN, NO, OO, SO) or two monodentate ligands and X is a monodentate moiety, often a leaving group e.g. Cl have been extensively studied as anticancer agents.⁶ These half-sandwich “piano-stool” complexes offer great scope for design, with the potential to vary each of the building blocks to allow modifications of thermodynamic and kinetic parameters. Indeed, it has been found that increasing the size of the coordinated arene increases their activity in the human ovarian cancer cell line. Changing the chelating ligand in these ruthenium arene complexes also appears to have an enormous effect on their kinetics and even changes their nucleobase selectivity.⁷

Synthesis and antiproliferative activity of Ru(II)(η⁶-arene) compounds carrying bioactive flavonol ligands have been reported by Hartinger et.al (A).⁸ Wei Su et al have described the DNA binding property and anticancer activity of ketone N4 substituted thiosemicarbazones and their ruthenium(II) arene complexes.⁹ A series of ruthenium(II) arene complexes with the 4-(biphenyl-4-carbonyl)-3-methyl-1-phenyl-5-pyrazolonate ligand, and related 1,3,5-triaza-7-phosphaadamantane (PTA) derivatives, have been reported along with their anticancer activity with low IC₅₀ value (B).¹⁰ Further, Dyson and his co-workers have reported the ruthenium(II)–arene complexes with a perfluoroalkyl-modified ligands displays remarkable in vitro cancer cell selectivity (C).¹¹ Recently, inhibitory activity of ruthenium(II) arene complexes of 2-phenylimidazole[4,5f][1,10]phenanthroline against the migration and invasion of MDA-MB-231 breast cancer cells have been investigated (D) (Figure 1).¹²
Figure 1. Reported ruthenium(II) arene anticancer drugs.

In recent years, much attention was given to compounds with pharmacophore hydrazone moieties due to the identification of several hydrazone lead compounds showing antiproliferative activity\(^1\) and antitumor activity\(^2\). It has been found from the literature that only a few reports are available on synthesis, characterisation and cytotoxicity of ruthenium(II) complexes containing hydrazone ligands\(^3\). Nevertheless, it should be pointed out that, as far as we know, the biological properties of arene ruthenium complexes bearing aroylhyrazones have not been studied so far. Therefore, in this study, we have combined ruthenium unit with a benzhydrazone ligand to generate a series of organometallic compounds with significant anticancer activity, taking advantage of the synthetic versatility of hydrazone derivatives and the promising biological activity.

We describe here, the synthesis and characterization of Ru(II) arene complexes containing bidentate indole-3-carboxaldehyde benzhydrazone ligands and chlorine. All the synthesized complexes have been characterized by elemental analysis, IR, UV-vis and NMR and ESI-MS spectroscopy techniques. The molecular structures of the complex 3 and 6 are confirmed through single crystal X-ray diffraction. The \textit{in vitro} cytotoxicity of the complexes 1-6 against HeLa, MDA-MB-231, Hep G2 and NIH 3T3 were screened by MTT assay. The morphological changes were investigated using various apoptosis assays (AO-EB staining, Hoechst staining, flow cytometry technique and comet assay). Further, the apoptosis pathway was confirmed by change in the mitochondrial membrane potential and western blot analysis.
Experimental Section

Methods and Instrumentation

The microanalysis of carbon, hydrogen, nitrogen and sulphur were recorded by an analytical function testing Vario EL III CHNS elemental analyser at the sophisticated Test and Instrumentation Centre (STIC), Cochin University, Kochi. Melting points were recorded with a Boetius micro-heating table and are corrected. Thermal measurements (TGA/DTA) were carried out on Perkin Elmer Thermal Analyzer in nitrogen atmosphere with a heating rate of 10 °C min⁻¹. FT-IR spectra were recorded in KBr pellets with JASCO 400 plus spectrometer. Electronic spectra in chloroform solution were recorded with a CARY 300 Bio UV-visible Varian spectrometer. ¹H NMR and ¹³C-NMR were spectra were recorded on a Bruker 400 MHz instrument using tetramethylsilane (TMS) as an internal reference. A Micro mass Quattro II triple quadrupole mass spectrometer was employed for electrospray ionization mass spectrometry (ESI-MS). The theoretical calculations were performed using the IsoPro software.¹⁶

Materials

The starting materials [(η⁶-C₆H₆)RuCl₂]₂ and (η⁶-p-cymene)RuCl₂]₂ were prepared according to literature methods.¹⁷

Procedure for the preparation of indole-3-carboxaldehyde benzhydrazones ligands

The ligands L1-L3 were prepared according to literature methods.¹⁸ A mixture of 4-substituted benzhydrazide (R=H, Cl or OMe derivatives) (1 mmol) and indole-3-carboxaldehyde (1 mmol) in ethanol (10 mL) containing a drop of glacial acetic acid was refluxed for 30 min. The separated solid was filtered and dried in air. Ligands were further purified by recrystallisation from methanol. Yield: 67-92%.

Procedure for the synthesis of ruthenium(II) arene benzhydrazone complexes

A mixture containing starting [(η⁶-arene)RuCl₂]₂ (arene-benzene or p-cymene) (0.05 mmol), indole-3-carboxaldehyde benzhydrazone ligand (0.1 mmol) and triethylamine (0.3 mL) in benzene (20 mL) was added and the resultant mixture was stirred at room temperature for 2 h. The orange brown precipitate was filtered, washed with hexane and dried in vacuo. The reaction progress was monitored through thin layer chromatography.

[Ru(η⁶-C₆H₆)(Cl)(L1)] (1): Brown solid. Yield = 0.160 g (68%); M.p.: 180⁰C (with decomposition); Calculated: C₂₂H₁₆ClN₃ORu: C, 55.40; H, 3.80; N, 8.81 %. Found: C, 55.37;
H, 3.79; N, 8.82 %. IR (KBr, cm\(^{-1}\)): 1539 \(v_{(C=N-N=C)}\), 1490 \(v_{(N-C-O)}\), 1369 \(v_{(C-O)}\). UV–Vis (CH\(_3\)CN, \(\lambda_{\text{max}}/\text{nm} \equiv \text{edm}^3 \text{ mol}^{-1} \text{ cm}^{-1}\)): 418 (1143), 273 (6371), 227 (14,757). \(^1\text{H} NMR\) (400 MHz, CDCl\(_3\)) (\(\delta\) ppm): 11.55 (br, 1H, indole N-H), 9.24 (s, 1H, HC=N), 7.08–7.98 (m, 10H, aromatic), 5.72 (s, 6H, CH-benzene). \(^13\text{C} NMR\) (400 MHz, CDCl\(_3\)) (\(\delta\) ppm) 164.15, 131.23, 129.73, 129.52, 129.20, 128.50, 127.50, 127.18, 125.05, 123.45, 122.45, 117.10, 116.82, 87.94 ppm. ESI-MS: displays a peak at m/z 441.56 (M - Cl\(^+\)) (calcd m/z 442.05).

**[Ru(\(\eta^5\)-C\(_6\)H\(_6\))(Cl)(L2)]\) (2):** Brown solid. Yield = 0.0933 g (69 %); M.p.: 172\(^0\)C (with decomposition); Calculated: C\(_{22}\)H\(_17\)Cl\(_2\)N\(_3\)ORu: C, 51.67; H, 3.35; N, 8.22 %. Found: C, 51.68; H, 3.36; N, 8.20 %. IR (KBr, cm\(^{-1}\)): 1531 \(v_{(C=N-N=C)}\), 1487 \(v_{(N-C-O)}\), 1378 \(v_{(C-O)}\). UV–Vis (CH\(_3\)CN, \(\lambda_{\text{max}}/\text{nm} \equiv \text{edm}^3 \text{ mol}^{-1} \text{ cm}^{-1}\)): 419 (1044), 269 (4977), 233 (10,051). \(^1\text{H} NMR\) (400 MHz, CDCl\(_3\)) (\(\delta\) ppm): 11.45 (br, 1H, indole N-H), 9.35 (s, 1H, HC=N), 6.78–7.92 (m, 9H, aromatic), 5.72 (s, 6H, CH-benzene). \(^13\text{C} NMR\) (400 MHz, CDCl\(_3\)) (\(\delta\) ppm) 162.73, 159.67, 130.61, 130.43, 129.30, 128.17, 114.54, 114.12, 88.57 ppm. ESI-MS: displays a peak at m/z 475.97 (M - Cl\(^+\)) (calcd m/z 476.01).

**[Ru(\(\eta^5\)-C\(_6\)H\(_6\))(Cl)(L3)]\) (3):** Orange brown solid. Yield = 0.268 g (92 %); M.p.: 186\(^0\)C (with decomposition); Calculated C\(_{23}\)H\(_{20}\)ClN\(_3\)O\(_2\)Ru: C, 54.49; H, 3.98; N, 8.29 %. Found: C, 54.48; H, 4.00; N, 8.29 %. IR (KBr, cm\(^{-1}\)): 1530 \(v_{(C=N-N=C)}\), 1486 \(v_{(N-C-O)}\), 1376 \(v_{(C-O)}\). UV–Vis (CH\(_3\)CN, \(\lambda_{\text{max}}/\text{nm} \equiv \text{edm}^3 \text{ mol}^{-1} \text{ cm}^{-1}\)): 429 (1496), 268 (4904), 236 (10,242). \(^1\text{H} NMR\) (400 MHz, CDCl\(_3\)) (\(\delta\) ppm): 11.45 (br, 1H, indole N-H), 9.36 (s, 1H, HC=N), 6.78–8.02 (m, 9H, aromatic), 5.72 (s, 6H, CH-benzene), 3.86 (s, 3H, OCH\(_3\)). \(^13\text{C} NMR\) (400 MHz, CDCl\(_3\)) (\(\delta\) ppm) 163.98, 136.49, 131.52, 128.90, 128.42, 126.99, 124.97, 124.97, 117.55, 117.00, 114.27, 114.27, 105.82, 88.94, 56.08 ppm. ESI-MS: displays a peak at m/z 471.99 (M - Cl\(^+\)) (calcd m/z 472.06). Single crystals suitable for X-ray diffraction were obtained by recrystallisation in DCM and methanol solution.

**[Ru(\(\eta^6\)-p-cymene)(Cl)(L1)]\) (4):** Orange-brown solid. Yield = 0.240 g (80 %); M.p.: 168\(^0\)C (with decomposition); Calculated: C\(_{26}\)H\(_{26}\)ClN\(_3\)ORu: C, 58.59; H, 4.92; N, 7.88 % Found: C, 58.59; H, 4.97; N, 7.85 %. IR (KBr, cm\(^{-1}\)): 1528 \(v_{(C=N-N=C)}\), 1486 \(v_{(N-C-O)}\), 1371 \(v_{(C-O)}\). UV–Vis (CH\(_3\)CN, \(\lambda_{\text{max}}/\text{nm} \equiv \text{edm}^3 \text{ mol}^{-1} \text{ cm}^{-1}\)): 431 (1044), 266 (4941), 228 (11,908). \(^1\text{H} NMR\) (400 MHz, CDCl\(_3\)) (\(\delta\) ppm): 11.86 (br, 1H, indole N-H), 9.29 (s, 1H, HC=N), 6.95–8.33 (m, 10H, aromatic), 5.58 (d, 1H, p-cym-H), 5.43 (d, 1H, p-cym-H), 5.40 (d, 1H, p-cym-H), 5.32 (d, 1H, p-cym-H), 2.85 (m, 1H, p-cym CH(CH\(_3\))\(_2\)), 2.31 (s, 3H, p-cym CH\(_3\)), 1.28 (d, 3H, p-cym CH(CH\(_3\))\(_2\)), 1.23 (d, 3H, p-cym CH(CH\(_3\))\(_2\)). \(^13\text{C} NMR\) (400 MHz, CDCl\(_3\)) (\(\delta\) ppm) 164.63, 151.89, 147.82, 146.37, 131.61, 131.06, 129.65, 129.38, 128.57, 127.37, 125.25, 123.62,
was solved with direct method using SIR97. Data were collected at 293K. The structure was refined by full matrix least-squares.

X-ray crystallography

Single crystals of [Ru(η⁶-C₆H₆)(Cl)(L3)] (3) and [Ru(η⁶-p-cymene)(Cl)(L3)] (6) were grown by slow evaporation of dichloromethane-methanol solution at room temperature. A single crystal of suitable size was covered with Paratone oil, mounted on the top of a glass fiber, and transferred to a Bruker AXS Kappa APEX II single crystal X-ray diffractometer using monochromated MoKα radiation (λ=0.71073). Data were collected at 293K. The structure was solved with direct method using SIR-97 and was refined by full matrix least-squares.
method on F2 with SHELXL-97. Non-hydrogen atoms were refined with anisotropy thermal parameters. All hydrogen atoms were geometrically fixed and collected to refine using a riding model. Frame integration and data reduction were performed using the Bruker SAINT Plus (Version 7.06a) software. The multi scan absorption corrections were applied to the data using SADABS software. Figure 1 was drawn with ORTEP and the structural data deposited at The Cambridge Crystallographic Data Centre: CCDC 1499166 and 1498893.

**Stability studies**

The stabilities of complexes 1–6 were checked by recording the UV-visible spectrum of them by dissolving in a minimum amount of 1% DMSO, and then diluted with PBS buffer. The hydrolysis profiles of these complexes were recorded by monitoring the electronic spectra for the resulting mixture over 24 h.

**Partition coefficients determination**

The hydrophobicity values of the complexes 1-6 were measured by the “Shake flask” method in octanol - water phase partitions as reported earlier. Complexes 1-6 (1 mg/mL) were dissolved in a mixture of water and n-octanol (2, 4, 6, 8, 10 μg/mL) followed by shaking for 1 hour. The mixture was allowed to settle over a period of 30 minutes and the resulting two phases were collected separately without cross contamination of one solvent layer into another. The concentration of the complexes in each phase was determined by UV-Vis absorption spectroscopy at room temperature. The results are given as the mean values obtained from three independent experiments. The sample solution concentration was used to calculate log P. Partition coefficients for 1-6 were calculated using the equation log P = log[(1-6)oct/(1-6)aq].

**Cell culture and inhibition of cell growth.**

**Cell culture.** HeLa (human cervical cancer cell line), MDA-MB-231 (Triple negative breast carcinoma), Hep G2 (human liver carcinoma cell line) and NIH 3T3 (noncancerous cell, mouse embryonic fibroblast) were obtained from the National Centre for Cell Science (NCCS), Pune. These cell lines were cultured as a monolayer in RPMI-1640 medium (Biochrom AG, Berlin, Germany), supplemented with 10% fetal bovine serum (Sigma-Aldrich, St. Louis, MO, USA) and with 100 U mL\(^{-1}\) penicillin and 100 μg mL\(^{-1}\) streptomycin as antibiotics (Himedia, Mumbai, India), at 37 °C in a humidified atmosphere of 5% CO\(_2\) in a CO\(_2\) incubator (Heraeus, Hanau, Germany).
Inhibition of cell growth

The IC\textsubscript{50} values, which are the concentrations of the tested compounds that inhibit
50\% of cell growth, were determined using a 3-(4,5-dimethyl thiazol-2-yl)-2,5-
diphenyl tetrazolium bromide (MTT) assay. Cells were plated in their growth medium
at a density of 5000 cells per well in 96 flat bottomed well plates. After 24 h plating,
the benzhydrazone ligands and Ru(II) arene benzhydrazone complexes 1-6 were added
at different concentrations (1-250 μM) for 24 h to study the dose dependent cytotoxic
effect. To each well, 20 μL of 5 mg mL\textsuperscript{-1} MTT in phosphate-buffer (PBS) was added.
The plates were wrapped with aluminium foil and incubated for 4 h at 37 °C. The
purple formazan product was dissolved by addition of 100 μL of 100% DMSO to each
well. The quantity of formazan formed gave a measure of the number of viable cells.
HeLa, MDA-MB-231 and Hep G2 were used for the MTT assay. The absorbance was
monitored at 570 nm (measurement) and 630 nm (reference) using a 96 well plate
reader (Bio-Rad, Hercules, CA, USA). Data were collected for four replicates each
and used to calculate the respective means. The percentage of inhibition was
calculated, from this data, using the formula: Percentage inhibition = 100 x \{Mean OD
of untreated cells (control) – Mean OD of treated cells\} /\{Mean OD of untreated cells
(control)\}. The IC\textsubscript{50} value was determined as the complex concentration that is
required to reduce the absorbance to half that of the control.

Acridine orange and ethidium bromide staining experiment

The changes in chromatin organization in MDA-MB-231 cells after treatment with
IC\textsubscript{50} concentration of the complexes 3 and 6 by using acridine orange (AO) and
ethidium bromide (EB). Briefly, about 5 x 10\textsuperscript{5} cells were allowed to adhere overnight
on a coverslip placed in each well of a 12-well plate. The cells were allowed to recover
for 1 h, washed thrice with DPBS, stained with an AO and EB mixture (1:1, 10 μM)
for 15 min, and observed with epifluorescence microscope (Carl Zeiss, Germany).

Hoechst 33258 staining method

Hoechst 33258 staining was done using the method described earlier with slight
modifications. 5 x 10\textsuperscript{5} MDA-MB-231 cells were treated with IC\textsubscript{50} concentration of
the complexes 3 and 6 for 24 h in a 6-well culture plate and were fixed with 4%
paraformaldehyde followed by permeabilization with 0.1% Triton X-100. Cells were
then stained with 50 μg mL⁻¹ Hoechst 33258 for 30 min at room temperature. The cells undergoing apoptosis, represented by the morphological changes of apoptotic nuclei, were observed and imaged by epifluorescence microscope (Carl Zeiss, Germany).

**Apoptosis evaluation - Flow cytometry**

The MDA-MB-231 cells were grown in a 6-well culture plate and exposed to IC₅₀ concentrations of complexes 3 and 6 for 24 h. The Annexin V-FITC kit uses annexin V conjugated with fluorescein isothiocyanate (FITC) to label phosphatidylserine sites on the membrane surface of apoptotic cells. Briefly the cells were trypsinised and washed with Annexin binding buffer and incubated with Annexin V-FITC and PI for 30 minutes and immediately analysed using flow cytometer FACS Aria-II. The results were analysed using DIVA software and percentage positive cells were calculated.

**Cellular DNA damage by the comet assay**

DNA damage was quantified by means of the comet assay as described. Assays were performed under red light at 4 °C. Cells used for the comet assay were sampled from a monolayer during the growing phase, 24 h after seeding. MDA-MB-231 cells were treated with the complexes 3 and 6 at IC₅₀ concentration, and cells were harvested by a trypsinization process at 24 h. A total of 200 μL of 1% normal agarose in PBS at 65 °C was dropped gently onto a fully frosted microslide, covered immediately with a coverslip, and placed over a frozen ice pack for about 5 min. The coverslip was removed after the gel had set. The cell suspension from one fraction was mixed with 1% low-melting agarose at 37 °C in a 1:3 ratio. A total of 100 μL of this mixture was applied quickly on top of the gel, coated over the microslide, and allowed to set as before. A third coating of 100 μL of 1% low-melting agarose was placed on the gel containing the cell suspension and allowed to set. Similarly, slides were prepared (in duplicate) for each cell fraction. After solidification of the agarose, the coverslips were removed, and the slides were immersed in an ice-cold lysis solution (2.5 M NaCl, 100 mM Na₂EDTA, 10 mM Tris, NaOH; pH 10, 0.1% Triton X-100) and placed in a refrigerator at 4 °C for 16 h. All of the above operations were performed in low-lighting conditions in order to avoid additional DNA damage. Slides, after removal from the lysis solution, were placed horizontally in an electrophoresis tank. The reservoirs were filled with an electrophoresis buffer (300 mM NaOH and 1 mM
Na<sub>2</sub>EDTA, pH > 13) until the slides were just immersed in it. The slides were allowed to stand in the buffer for about 20 min (to allow DNA unwinding), after which electrophoresis was carried out at 0.8 v cm<sup>-1</sup> for 15 min. After electrophoresis, the slides were removed, washed thrice in a neutralization buffer (0.4 M Tris, pH 7.5), and gently dabbed to dry. Nuclear DNA was stained with 20 μL of EB (50 μg mL<sup>-1</sup>). Photographs were taken using an epifluorescence microscope (Carl Zeiss).

**Mitochondrial membrane potential Assay**

Mitochondrial membrane potential, Δψ<sub>m</sub> is an important parameter of mitochondrial function used as an indicator of cell health. MDA-MB-231 cells treated overnight with IC<sub>50</sub> concentration of the complexes 3 and 6 in 6-well plates were incubated for 1 h with 2 μg mL<sup>-1</sup> of JC−1 in the culture medium. The adherent cell layer was then washed three times with PBS and dislodged with 250 μL of trypsin–EDTA. Cells were collected in PBS/2% bovine serum albumin (BSA), washed twice by centrifugation, resuspended in 0.3 mL of PBS/2% BSA, mixed gently, and examined in the fluorescent microscope (Carl Zeiss, Jena, Germany).

**Western blot analysis**

For Western blotting, MDA-MB-231 cells were treated with the complexes 3 and 6 at IC<sub>50</sub> concentration for 24 h, and appropriate amounts of cell lysates (25 μg protein) were resolved over 10% Tris–glycine polyacrylamide gel, and then transferred onto the PVDF membrane. The blots were blocked using 5% non-fat dry milk and probed using p53, Bcl-2 and Bax primary monoclonal antibodies in blocking buffer overnight at 4 °C. The membrane was then incubated with appropriate secondary antibody-horseradish peroxidase conjugate (Amersham Life Sciences Inc., IL, USA), followed by detection using chemiluminescence ECL kit (Amersham Life Sciences Inc., IL, USA). To ensure equal loading of protein, the membrane was stripped and reprobed with anti-β-actin antibody (Sigma Aldrich, USA).

**Results and Discussion**

**Synthesis of ruthenium(II) arrene benzhydrazone complexes**

The hydrazone ligand derivatives were conveniently prepared in an excellent yield by the condensation of indole-3-carboxaldehyde with 4-substituted benzhydrazides (H, Cl and...
OMe derivatives) in an equimolar ratio. These ligands were allowed to react with the ruthenium(II) arene precursor [[(η$_6$-arene)RuCl$_2$]$_2$ (arene-benzene or p-cymene) in a 2:1 molar ratio in the presence of triethylamine as the base and the new complexes of the general formula, [(η$_6$-arene)Ru(L)Cl] (arene-benzene or p-cymene; L-substituted indole-3-carboxaldehyde benzhydrazone derivatives) (Scheme 1) were obtained in high yields. The addition of triethylamine to the reaction mixture was used to remove a proton from the imidolate oxygen and to facilitate the coordination of the imidolate oxygen to the ruthenium(II) ion. All complexes are air-stable and are highly soluble in most organic solvents. The analytical data of all the ruthenium(II) arene benzhydrazone complexes are in good agreement with the molecular formula proposed.

![Scheme 1](image)

Scheme 1. Synthesis of ruthenium(II) arene indole-3-carboxaldehyde benzhydrazone complexes

Characterization of the complexes

The IR spectra of the free ligands displayed a medium to strong band in the region of 3180-3196 cm$^{-1}$ which is characteristic of the N-H functional group. The free ligands also displayed $v$C=N and $v$C=O absorptions in the region of 1548-1576 cm$^{-1}$ and 1610-1653 cm$^{-1}$ respectively, which indicate that the ligands exist in the amide form in the solid state. Bands that are due to $v$N-H and $v$C=O stretching vibrations were not observed with the complexes, which indicates that the ligands underwent tautomerization and subsequent coordination of the imidolate enolate form during complexation. Coordination of the ligand to the ruthenium(II) ion through an azomethine nitrogen is expected to reduce the electron density in the azomethine link and thus lower the absorption frequency upon complexation 1528-1539 cm$^{-1}$ which indicates the coordination of azomethine nitrogen to the ruthenium(II) ion. The band in the region of 1369-1378 cm$^{-1}$ is due to the imidolate oxygen, which is
coordinated to the metal. The IR spectra of all the complexes therefore confirm the mode of
coordination of the benzhydrazone ligand to the ruthenium(II) ion via the azomethine
nitrogen and imidolate oxygen.\textsuperscript{21}

The absorption spectra of the ruthenium(II) arene benzhydrazone complexes in
chloroform exhibited very intense band around 266-273 nm and 227-236 nm are assigned to
ligand-centered (LC) $\pi-\pi^*$ and $n-\pi^*$ transitions respectively. The lowest energy absorption
bands in the electronic spectra of the complexes in the visible region 410-431 nm are ascribed
to MLCT (metal to ligand charge transfer) transitions. Based on the pattern of the electronic
spectra of all the complexes an octahedral environment around the ruthenium(II) ion has been
proposed similar to that of the other octahedral ruthenium(II) arene complexes.\textsuperscript{22}

The $^1$H NMR spectra of all the complexes were recorded in CDCl$_3$ to confirm the
bonding of the benzoylhydrazone ligand to the ruthenium(II) ion. Multiplets observed in the
region $\delta$ 6.74-8.61 ppm in the complexes have been assigned to the aromatic protons of
benzhydrazone ligands. The signal due to the azomethine proton appears in the region $\delta$ 9.24-
9.49 ppm. The position of the azomethine signal in the complexes is slightly downfield in
comparison with that of the free ligand, suggesting deshielding of the azomethine proton due
to its coordination to ruthenium. The singlet due to the -NH proton of the free ligand in the
region $\delta$ 11.22-11.60 ppm is absent in the complex, further supporting enolisation and
coordination of the imidolate oxygen to the Ru(II) ion. Therefore, the $^1$H NMR spectra of the
complexes confirm the bidentate coordination mode of the benzhydrazone ligands to
ruthenium(II) ion. In all the complexes, the indole N-H protons are observed as singlets in
between $\delta$ 11.41-11.88 ppm. The cymene protons are appeared in the region of $\delta$ 5.32-5.62
ppm.\textsuperscript{23} In addition, the two isopropyl methyl protons of the p-cymene appeared as two
doublet in the region of $\delta$ 1.23-1.41 ppm and the methine protons comes in the region of $\delta$
2.31-3.10 ppm as septet. Further, the methyl group of the p-cymene comes as singlet around
the region of $\delta$ 2.31-2.34 ppm. Additionally methoxy protons are observed as singlet for
complexes 3 and 6 at $\delta$ 3.81-3.86 ppm. On the other hand, benzene arene protons displayed
an upfield shift relative to complex 4-6 in the region $\delta$ 5.72-5.73 ppm. (Figure S1, Supporting
Information). The $^{13}$C NMR of the Ru(II) arene complexes showed resonance in the expected
regions (Figure S2, Supplementary material) and the complex revealed a downfield shift of
the azomethine carbon relative to the free ligands indicating coordination of the azomethine
nitrogen to the metal centre.
Stability of Complexes (Time-Dependent Spectra).

Stability in solution is an important requirement for drug candidates. The stability of the most cytotoxic complexes 1-6, was studied by UV–Vis spectroscopy in a solution of 1% DMSO in PBS. All the ruthenium(II) arene benzhydrazone complexes showed characteristic peaks in the region of 200-800 nm and did not exhibit any significant changes during a 24-hour period. The absence of significant changes in the peak absorptions and spectral characteristics for tested complexes over time may suggest that no structural alternations occurred in buffer solution. The data for all studied complexes is presented in Figure S5, (Supporting Information). Further, the composition of the complexes has been studied by ESI-MS spectral studies. Mass spectrometric measurements carried out under positive ion ESI mode using acetonitrile as the solvent. In their positive ESI mass spectra 1-6 showed major peaks due to cationic fragment \([\eta^6\text{-arene}Ru(L)Cl]^+\) generated by loss of the Cl. The ESI spectra of complexes 1-6 display at m/z found (calcd): [441.56 (442.05) (1, M - Cl)], [475.97 (476.01) (2, M - Cl)], [471.99 (472.06) (3, M - Cl)], [497.62 (498.12) (4, M - Cl)], [531.21 (532.08) (5, M - HCl)] and [527.86 (528.13) (6, M - Cl)] respectively confirm the presence of monomeric entity in solution phase. The mass spectrometry results are in good agreement with the proposed molecular formulae of the complexes and suggest that chloro (Cl) group is labile and possibly replaced by targeted biomolecules. The experimentally observed and theoretically calculated isotopic distributions were in excellent agreement with each other as illustrated in and are shown in Figure S3 and S4, (Supporting Information). Further, the thermal stability of the synthesized ruthenium(II) arene complexes 3 and 6 was determined by thermogravimetric analysis (TGA) and differential thermal analysis (DTA) as shown in S6 (Supporting Information). The synthesized complex is stable up to 180 °C. The results are in good agreement with the formulae suggested from the analytical data.

X-ray crystallographic studies

Attempts were made to grow single crystals for all the complexes to confirm the coordination mode of the ligand to metal and geometry of the complex. However we could obtained single crystals for complexes \([\text{Ru}(\eta^6\text{-p-cymene})(\text{Cl})(L3)]\) (3) and \([\text{Ru}(\eta^6\text{-C}_6\text{H}_6)(\text{Cl})(L3)]\) (6). Crystals of 3 and 6 grew from slow diffusion of dichloromethane into methanol solutions and crystallized in the monoclinic system with P2(1)/n space group. The selected bond lengths and bond angles are given in Table 1 whereas crystallographic data and structural refinement parameters are gathered in Table S1 (Supporting Information).
ORTEP views of the molecules with the atom numbering are shown in Figures 2 and 3. The molecular structure of the complex 3 shows clearly that the benzhydrazine ligand coordinates in a bidentate manner to ruthenium ion via the azomethine nitrogen and imidolate oxygen in addition to one chlorine and one arene group. The complex adopts the commonly observed piano-stool geometry as reported in many half-sandwich arene ruthenium(II) complexes. In this case, the arene ring forms the seat of the piano-stool, while the bidentate benzhydrazine N, O and Cl ligands form the three legs of the stool. Therefore, ruthenium(II) ion is sitting in a NOCl ($\eta^6$-arene) coordination environment. The benzhydrazine ligand bind to the metal centre at N and O forming the five membered chelate ring with bite angle 76.18(12)$^\circ$ O(1)-Ru(1)-N(2) and 84.26(11)$^\circ$ N(2)-Ru(1)-Cl(1). The bond lengths of Ru(1)-N(2) and Ru(1)-O(1) are 2.069(3) Å and 2.069(3) Å respectively. The Ru-Cl bond length is found to be 2.4233(13) Å and the bond length is in agreement with other structurally characterized $p$-cymene ruthenium complexes. The ruthenium atom is $\pi$ bonded to the arene ring with an average Ru-C distance of 2.156(7) Å, whereas average C-C bond length in the arene ring is 1.425(8) Å with alternating short and long bonds. It was observed that the complex 6 also adopt similar geometrical environment as in the complex 3 with slight variation in the bond angles and bond distances. The crystal structures of 3 and 6 revealed the presence of extensive intermolecular hydrogen bonding interactions shown in (Figure S7, Supporting Information).

Table 1. Selected Bond Lengths (Å) and Angles ($^\circ$) in 3.H$_2$O and 6.

<table>
<thead>
<tr>
<th>distances/angles</th>
<th>3.H$_2$O</th>
<th>6</th>
</tr>
</thead>
<tbody>
<tr>
<td>Ru1-N2</td>
<td>2.069(3)</td>
<td>2.077(4)</td>
</tr>
<tr>
<td>Ru1-O1</td>
<td>2.069(3)</td>
<td>2.065(4)</td>
</tr>
<tr>
<td>Ru1-Cl1</td>
<td>2.4233(13)</td>
<td>2.4060(16)</td>
</tr>
<tr>
<td>Ru1-C22</td>
<td>2.156(7)</td>
<td>2.153(5)</td>
</tr>
<tr>
<td>N1-N2</td>
<td>1.391(5)</td>
<td>1.389(6)</td>
</tr>
<tr>
<td>N2-C7</td>
<td>1.326(6)</td>
<td>1.326(6)</td>
</tr>
<tr>
<td>O2-C9</td>
<td>1.289(5)</td>
<td>1.289(6)</td>
</tr>
<tr>
<td>O1-Ru1-N2</td>
<td>76.18(12)</td>
<td>75.71(16)</td>
</tr>
<tr>
<td>N2-Ru1-Cl1</td>
<td>84.26(11)</td>
<td>83.85(13)</td>
</tr>
<tr>
<td>N2-N1-Ru1</td>
<td>116.0(2)</td>
<td>115.7(3)</td>
</tr>
<tr>
<td>C7-O1-Ru1</td>
<td>112.8(2)</td>
<td>112.8(3)</td>
</tr>
<tr>
<td>C7-N2-N1</td>
<td>110.5(3)</td>
<td>110.6(4)</td>
</tr>
<tr>
<td>C19-Ru1-Cl1</td>
<td>106.2(3)</td>
<td>104.81(16)</td>
</tr>
<tr>
<td>O1-Ru1-Cl1</td>
<td>84.46(10)</td>
<td>86.90(12)</td>
</tr>
</tbody>
</table>
Figure 2. ORTEP drawing of complex 3.H₂O at 30% probability level.

Figure 3. ORTEP drawing of complex 6 at 30% probability level.

Partition coefficients determination

Lipophilicity is an important factor for the cellular accumulation and oral bioavailability of drugs. It is often expressed as the n-octanol/water partition coefficient (log P), which is also a central parameter in many in silico medicinal chemistry approaches, such as the determination of the drug likeliness of a new drug. This was investigated by the partition coefficient, P a parameter, which indicates the hydrophobic character of molecules and their ability to cross lipid bilayers. The calculated log P values for complexes 1-6 are 2.59, 2.72, 2.48, 2.23, 2.35 and 1.99 respectively. It has been observed that complex 6 with a p-cymene group shows higher potency than the rest of the complexes (Table 2).
**In Vitro antiproliferative activity**

All the ruthenium complexes and the free benzhydrazone ligands evaluated for their cytotoxic activity against HeLa, MDA-MB-231 and Hep-G2 along with NIH 3T3 cell lines by using colorimetric assay (MTT assay) that measures mitochondrial dehydrogenase activity as an indication of cell viability. The effects of the ruthenium(II) arene complexes to arrest the proliferation of cancer cells were evaluated after an exposure of 24 h. It is to be noted that the ligands did not show any inhibition of the cell growth even up to 100 μM and clearly indicates chelation of the ligand with metal ion is responsible for the observed cytotoxicity properties of the complexes. The results of MTT assays revealed that complexes showed notable activity against the cell lines HeLa, MDA-MB-231 and Hep-G2 with respect to IC$_{50}$ values (Table 2). From the IC$_{50}$ values obtained it was inferred that complexes 3, 4 and 6 are highly active against all the cell lines with very low IC$_{50}$ values than that of the well-known anticancer drug cisplatin. In addition, the *in vitro* cytotoxic activity studies of the complexes against the mouse embryonic fibroblast cell line NIH 3T3 (normal cells) was undertaken and the IC$_{50}$ values are above 215 μM, which confirmed that the complexes are very specific on cancer cells.

These ruthenium(II) arene benzhydrazone complexes 1-6 possess significant cytotoxicity over the ligands may be due to the presence of extended π conjugation resulting from the chelation of Ru(II) ion with the ligand. Further, the observed higher activity of the complexes 4 and 6 is correlated to the nature of the chelating benzoylhydrazone ligand and arene moiety. Further, the observed higher activity of the complexes 3 and 6 is correlated to the nature of the chelating benzhydrazone ligand and arene moiety. In the complexes 3 and 6, the presence of electron donating methoxy substituent at phenyl ring of the ligand increases the lipophilic character of the metal complex, which favours its permeation through the lipid layer of a cell membrane.

On the other hand, the arene groups also play an important role in the antitumor activity of these ruthenium complexes. It has been observed that complex 6 with a p-cymene group show higher potency than those with a benzene group in complex 3, which may be attributed to the stronger hydrophobic interactions between the Ru (II)-cymene complex and the biomolecular targets as evidenced by partition coefficient value. Complex 6 shows high cytotoxic activity with very low IC$_{50}$ values of 11.4 ± 0.7, 4.1 ± 0.4 and 9.1 ± 0.3 μM toward HeLa, MDA-MB-231 and Hep-G2. Further, the IC$_{50}$ values are much better than those previously reported for other Ru(II) arenylazo, 2-thiosalicylic acid, phenanthroimidazole
or polypyridyl complexes. The excellent results suggest to investigate the underlying mechanism accounting for the antiproliferative action of these ruthenium arene benzhydrazone complexes.

**Table 2.** Cytotoxicity (IC$_{50}$, µM) of ligand and complexes 1-6. (n.e.: no effect) and calculated partition coefficients (log $P$).

<table>
<thead>
<tr>
<th>Complex</th>
<th>IC$_{50}$ values (µM)</th>
<th>log $P$</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>HeLa</td>
<td>MDA-MB-231</td>
</tr>
<tr>
<td>Complex 1</td>
<td>20.8 ± 0.2</td>
<td>18.2 ± 0.8</td>
</tr>
<tr>
<td>Complex 2</td>
<td>25.9 ± 0.8</td>
<td>19.9 ± 0.1</td>
</tr>
<tr>
<td>Complex 3</td>
<td>19.4 ± 0.3</td>
<td>15.3 ± 0.3</td>
</tr>
<tr>
<td>Complex 4</td>
<td>13.6 ± 0.4</td>
<td>11.2 ± 0.3</td>
</tr>
<tr>
<td>Complex 5</td>
<td>17.9 ± 0.3</td>
<td>12.8 ± 0.2</td>
</tr>
<tr>
<td>Complex 6</td>
<td>11.4 ± 0.7</td>
<td>4.1 ± 0.4</td>
</tr>
<tr>
<td></td>
<td>L1</td>
<td>n.e.</td>
</tr>
<tr>
<td></td>
<td>L2</td>
<td>n.e.</td>
</tr>
<tr>
<td></td>
<td>L3</td>
<td>n.e.</td>
</tr>
<tr>
<td></td>
<td>Cisplatin</td>
<td>19.2 ± 1.1</td>
</tr>
</tbody>
</table>

The ligands L1-L3 were added at different concentrations (1-250 µM) for 24 h.

**AO-EH and Hoechst staining assays**

Acridine orange and ethidium bromide (AO and EB) dual staining followed by fluorescence microscopy revealed apoptosis from the perspective of fluorescence emission. Apoptosis is characterized by cell shrinkage, blebbing of the plasma membrane and chromatin condensation. To identify apoptosis, at a basic level, we adopted AO-EB staining to visualize and quantify the number of viable and apoptic cells. According to the difference in membrane integrity between necrotic and apoptosis, AO can pass through cell membrane, but EB cannot. After the treatment of MDA-MB-231 cells with complexes 3 and 6 for 24 h at IC$_{50}$ concentration, the apoptotic effect is shown in Figure 4. The cells incubated with the complexes 3 and 6 for 2 h and irradiated with visible light showed significant reddish-orange emission characteristic of the apoptotic cells. In the control, the cells of MDA-MB-231 were stained bright green in spots. Additionally, complexes 3 and 6 treated MDA-MB-231
cells were stained with Hoechst 33258, apoptotic features such as nuclear shrinkage and chromatin condensation were also observed (Figure 5). Hence the results of AO-EB and Hoechst staining assays suggest that complexes 3 and 6 induce apoptosis in MDA-MB-231 cells.  

**Figure 4.** Morphological assessment of AO and EB of MDA-MB-231 cells treated with complexes 3 and 6 (IC$_{50}$ concentration) for 24 h. The scale bar 20 µm.

**Figure 5.** Morphological assessment of complexes 3 and 6 (IC$_{50}$ concentration) and MDA-MB-231 cells for 24 h. The scale bar 20 µm.

**Evaluation of apoptosis - Flow cytometry**

The potential to induce apoptosis in cancer cells by the addition of synthesized complexes can be quantitatively investigated by flow cytometry analysis by Annexin V protocol, with the help of Annexin V-FITC Apoptosis Detection Kit to perform double-staining with propidium iodide and Annexin V-FITC. Annexin V, a Ca$^{2+}$ dependent phospholipid-binding protein with a high affinity for the membrane phospholipid phosphatidylserine (PS), is quite helpful for identifying apoptotic cells with exposed PS. Propidium iodide is a standard flow cytometric viability probe used to distinguish viable from non-viable cells (Figure 6). The MDA-MB-231 cells were treated with the complexes 3 and 6 at IC$_{50}$ concentrations for 24 h. The cell death induced by the complexes follow a pathway from lower left quadrant to the upper right quadrant (Annexin V$^+$/PI$^+$) which represents cells undergoing apoptosis.
Figure 6. AnnexinV/propidium iodide assay of MDA-MB-231 cells treated by complexes 3 and 6 (IC$_{50}$ concentration) measured by flow cytometry.

Comet assay

The Comet Assay (single-cell gel electrophoresis) in an agarose gel matrix was used to study DNA fragmentation. The comet assay was performed with treated MDA-MB-231 cancer cells with IC$_{50}$ concentration of complexes 3 and 6, large and well-rounded comets are observed while the control cells fail to show a comet like appearance (Figure 7). The comet score for complexes 3 and 6 shows significant number of nucleoids with larger comet tails, indicative of higher levels of DNA single-strand breaks.$^{31}$

Figure 7. Comet assay of staining of EB control (untreated) treated with complexes 3 and 6 (IC$_{50}$ concentration) for 24 h. The scale bar 40 µm.

Mitochondrial membrane potential detection

Mitochondria act as a point of integration for apoptotic signals originating from both extrinsic and intrinsic apoptotic pathways. Mitochondria play important roles in apoptosis through the release of proapoptotic factors such as cytochrome $c$ and other apoptosis-inducing factors. The changes in mitochondrial membrane potential were detected using the fluorescent probe JC-1. It exhibits potential-dependent
accumulation in mitochondria, indicated by a fluorescence emission shift from red
(~590 nm) to green (~525 nm). As shown in Fig, in the control, JC-1 emits red
fluorescence. When the MDA-MB-231 cells were treated with the complexes, JC-1
displays a green fluorescence. The changes from red to green fluorescence indicate the
decrease of mitochondrial membrane potential (Figure 8). These results suggest that
complexes 3 and 6 can induce the decrease of mitochondrial membrane potential.32

Figure 8. MDA-MB-231 cells were treated with complexes 3 and 6 (IC50 concentration) for
24 h. The scale bar 20 µm

Western blot analysis
To reveal the underlying mechanism behind the antiproliferative activity of Ru(II)
benzhydrazone complexes, Western blot technique has been employed. It is
established that apoptotic proteins like p53, Bax and anti-apoptotic protein Bcl-2 play a
pivotal role during the induction of apoptosis. The expression level of p53, Bax and
Bcl-2 proteins were analyzed in the 3 and 6 treated MDA-MB-231 cells and control
cells. It is observed that the expression level of the Bcl-2 protein decreases suggesting
that apoptosis by 3 and 6 could be mediated through the downregulation of the
antiapoptotic protein Bcl-2. The p53 and Bax proteins level in MDA-MB-231 cancer
cell lines is remarkably increased upon treatment with the complexes revealing that the
complexes induce apoptosis (Figure 9). Hence, the upregulation of proapoptotic
protein Bax, p53 and the downregulation of antiapoptotic protein Bcl-2 caused by
complexes 3 and 6 could possibly activate mitochondria-mediated apoptosis.33
Figure 9. Western blot of p53, Bax and Bcl–2 proteins in MDA-MB-231 cells. Lane-1 control, lanes-2 and 3 treated with the complexes 3 and 6 (IC₅₀ concentration). β–Actin as a loading control.

Conclusions

An easy route of synthesis of six new ruthenium(II) arene indole-3-carboxaldehyde benzhydrazone has been described for the first time. The characterization of the complexes (1-6) was accomplished by analytical and spectral methods (IR, UV–vis, $^1$H and $^{13}$C NMR and ESI-MS). X-ray diffraction study reveals that the benzhydrazone ligand coordinated to ruthenium via azomethine nitrogen and imidolate oxygen and adopts the familiar pseudo-octahedral “piano-stool” geometry. Interestingly, the cytotoxic activities of complex 6 against the tested cancer cell lines were significantly superior to that of the well-known anticancer drug cisplatin and the observed high cytotoxicity is correlated with nature of the substituent of the ligand and arene moiety. Furthermore, fluorescence staining techniques and flow cytometry using the annexin-V assay revealed that complexes 3 and 6 induce apoptosis in MDA-MB-231 cancer cells. Further alkaline comet assay confirms the single-strand break of DNA. The results of mitochondrial membrane potential and Western blot analysis demonstrated that the complexes with potent antiproliferative activity are able to induce mitochondria-mediated apoptosis in human cancer cells. On the basis of the results, we suggest that ruthenium arene benzhydrazone complexes may be the best candidates for further evaluation as chemopreventive and chemotherapeutic agents for human cancers.
Acknowledgements

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Supporting Information

CCDC deposition No. 1499166 (3) and 1498893 (6) contain the supplementary crystallographic data for this paper. In addition, selected crystal data and structure refinement data and Figures contain the $^1$H and $^{13}$C NMR, ESI-MS, UV-vis spectrum and Intermolecular interaction diagrams of complexes 3 and 6 are provided.

References


16. (a) J. Fernandez-de-Cossio, *Analytical Chemistry*, 2010, **82**, 1759-1765; (b) IsoPro, version 3.1; Cornell University. A computer program written by Michael W. Senko that implements Yergey’s polynomial method running under Microsoft Windows.


A series of ruthenium(II) arene complexes have been synthesized and evaluated for their \textit{in vitro} anticancer activities. The complex exhibits promising anticancer activity in human cancer cells.