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SYNTHESIS, CHARACTERIZATION AND DNA BINDING ACTIVITY OF A POTENTIAL DNA INTERCALATOR

(Sintesis dan Pengikatan DNA oleh Sebatian Interkalasi DNA Yang Berpotensi)

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Abstract

A novel complex, $[Ru(dppz)_2(p\text{-MOPIP})]^{2+}$ (dppz = dipyrido-[3,2-a:20,30-c]phenazine, p-MOPIP = 2-(4-methoxyphenyl) imidazo[4,5-f][1,10]phenanthroline) has been synthesized and characterized by elemental analysis, 1H Nuclear Magnetic Resonance spectroscopy, mass spectrometry, Fourier Transform Infrared analysis, Ultra Violet visible and fluorescence spectroscopy. Herein, the complex was designed by adding p-MOPIP as an intercalating ligand and dppz as the ancillary ligand. The DNA binding properties of the complex with Calf Thymus DNA (CT-DNA) were investigated using spectroscopic methods. The UV-visible absorption band observed at 460 nm corresponded to the metal-to-ligand charge transfer (MLCT) while bands at 358 and 281 nm corresponded to intra-ligand (IL) π - π * transitions of the ligand scaffold in p-MOPIP and dppz. The intrinsic binding constant, K_b for this complex was $1.67 \times 10^6 \, \mathrm{M}^{-1}$ and this suggested that this complex, $[Ru(dppz)_2(p\text{-MOPIP})]^{2+}$ bound to DNA via the intercalative mode. Interestingly, the interaction of this complex with CT-DNA also had a 'molecular light switch' effect.

Keywords: ruthenium, DNA binding, molecular light switch

Abstrak

Satu kompleks baru, $[Ru(dppz)_2(p-MOPIP)]^{2+}$ (dppz = dipyrido-[3,2-a:20,30-c]fenazin, p-MOPIP = 2-(4-metoksifenil) imidazo[4,5-f][1,10]fenantrolina) telah dihasilkan dan dianalisis dengan menggunakan analisis elemen, spektroskopi resonans magnet nukleus (1 H NMR), spektrometri jisim, analisis inframerah tranformasi Fourier (FTIR), spektroskopi ultra-lembayung dan pendafluor. Penyediaan kompleks telah dilakukan dengan menambah p-MOPIP sebagai ligan interkalasi dan dppz sebagai ligan sisi. Ciri-ciri ikatan DNA bagi kompleks ini dengan Calf Thymus DNA (CT-DNA) telah dianalisis dengan menggunakan kaedah spektroskopik. Jalur pada 460 nm merujuk kepada caj metal-kepada-ligan (MLCT), manakala jalur pada 358 dan 281 nm dikaitkan dengan peralihan π - π * intra-ligan (IL) bagi ligan p-MOPIP dan dppz. Nilai pemalar ikatan, K_b ialah 1.67 x 10 6 M $^{-1}$ dan keputusan eksperimen mencadangkan pengikatan berlaku melalui mod interkalasi. Menariknya, kompleks ini juga menunjukkan kesan 'suis cahaya molekul'.

Kata kunci: ruthenium, pengikatan DNA, suis cahaya molekul

Introduction

Ruthenium, as other transition metals, generally have the electronic configuration where the 4d sub-shells are partially filled [1], allowing it to form different sorts of complexes that are suitable for various applications including as anticancer drugs, biosensors, in catalysis and others. Ruthenium complexes have properties that make them active biologically [2]. One of the properties is ligand exchange, where ruthenium(II) and ruthenium(III)

complexes having similar exchange kinetics to those of platinum(II) complexes [3], thus, are able to display similar biological effects as platinum(II) drugs. Additionally, ruthenium has a wide range of accessible oxidation states [4], for instance, ruthenium(II), ruthenium(III) or ruthenium(IV). The reduction and oxidation (redox) potential of metal complexes can be modified and exploited by varying the ligands to improve the effectiveness of ruthenium based drugs in the pharmaceutical industry [5]. Ruthenium(II) complexes with d⁶ configuration are predominantly are octahedral and diamagnetic [6]. Recently, researchers have been more focused on the ruthenium(II) polypyridyl complexes due to their photoluminescence which makes them stable and also suitable as DNA probes.

Previously, the development of classical ruthenium drugs was based on the capability of ruthenium to bind with DNA *via* some of the nitrogen atoms of nucleic bases. The interaction between transition metal complexes and nucleic acids were studied extensively in order to develop drugs that could react and bind to nucleic bases. The most important characteristics of ruthenium(II) complexes are that their intercalating ligand must possess extended and planar aromatic structures which is able to stack and insert between the base pairs of the DNA double helix. The intercalating ligand plays an important role in order to make the interaction with DNA happen.

The modification of polypyridyl ligands would allow fine tuning of the configuration and electronic structures of ruthenium(II) polypyridyl complexes therefore, varying the DNA binding properties of these complexes. Nowadays, the researchers are more interested in the complexes that contain one intercalating ligand and two ancillary ligands due to their stability, strong DNA affinity, photochemical properties and various other applications. Ruthenium complexes with stable, inert, coordinatively saturated and water soluble properties are the most ideally suited and extremely valuable as non-covalent probes of both structural and functional aspects of nucleic acid chemistry [7].

Thus, the objective of this study is to synthesis, characterization and evaluate the DNA binding activity of a potential DNA intercalator based on novel complex, $[Ru(dppz)_2(p-MOPIP)]^{2+}$.

Materials and Methods

Materials

All the chemicals and solvents in this experiment are at least of analytical grade and used as received from the suppliers unless stated otherwise. The reagents used without purification are as follows: ruthenium(III) chloride hydrate, 4-methoxybenzaldehyde, 1,10-phenanthroline, sodium bromide, sodium hydrogen carbonate, magnesium sulfate, ammonium acetate, *o*-phenylene diamine, lithium chloride (LiCl), potassium hexaflorophosphate (KPF₆) and tetra-*N*-butylammoniumchloride. The solvents are listed as follows: nitric acid, sulphuric acid, chloroform, methanol, glacial acetic acid, aqueous ammonia, dimethylformamide, toluene, acetonitrile and ethylene glycol.

Instrumentation

The elemental analysis of the complex was carried out by using Leco CHNS-932 Elemental Analyzer. The nuclear magnetic resonance (NMR) spectra was recorded at room temperature in deuterated acetonitrile (CD $_3$ CN) by using JEOL ECX500 FT NMR 500 MHz spectrometer with tetramethylsilane (TMS) as the internal standard for 1 H. Electrospray Ionization Mass Spectrometry (ESI-MS) of the complex was performed on a triple stage quadrupole Finnigan TSQ7000 mass spectrometer with a diverted inject valve where the spectra recorded were in the scanning range of 50 – 2200 m/z ratio. The Infrared absorption spectra were measured by using Perkin Elmer 100 series FT-IR Spectrometer in the range 4000 – 200 cm $^{-1}$. UV-Visible spectra were recorded over the range of 200 – 600 nm on Shimadzu H.U.V.1650 PC UV-Visible spectrophotometer using quartz cells of 10 mm path length. Emission spectra were recorded over the range 500 – 850 nm using an excitation wavelength of 460 nm on a Shimadzu RF-5301 PC spectrofluorometer at room temperature.

Preparation of the precursor ligand and complex

The precursor ligand, 1,10-phenanthroline-5,6-dione was prepared according to literature [8] and recrystallized in methanol. The precursor ligand was later used to prepare dipyrido[3,2-a:2',3'-c]phenazine (dppz) [9] and 2-(4-methoxylphenyl)imidazo[4,5-f][1,10]phenanthroline (p-MOPIP) [10] by following literature methods.

Synthesis of Ru(dppz)₂Cl₂

Ru(dppz)₂Cl₂ was prepared according to an established method [11]. A mixture of 0.2 g (1.0 mmol) RuCl₃.3H₂O, 0.55 g (2.0 mmol) dppz, 0.22 g of LiCl, and 30 mL of DMF was refluxed for 8 hours. The dark solution was stirred in 100 mL of chilled acetone and then kept in the freezer overnight. The crude product was collected by filtration and washed with water. The dried crude product was a fine dark purple powder. Yield: 0.623 g (80.1%), MS, m/z: 703 [M⁺- 2Cl]; ¹H NMR 500 MHz (CD₃CN, δ ppm): 9.50 (d, 4H), 9.17 (d, 4H), 8.38 (dd, 4H), 8.04 (dd, 4H) and 7.92 (d, 4H); IR (v, cm⁻¹):3057.49 (sp² C-H), 1631.00 (C=N), 1581.31 (C=C), 1073.58 (C-N) and 327.32 (Ru-Cl).

Synthesis of $[Ru(dppz)_2p-MOPIP_2]^{2+}$

A mixture of 0.034g (0.10 mmol) p-MOPIP, 0.08g (0.10 mmol) of $[Ru(dppz)_2Cl_2]^{2+}$ was dissolved in 10 ml of ethylene glycol and refluxed for 4 hours under nitrogen. Upon cooling, concentrated KPF₆ solution was added into the dark solution. Once the precipitation was complete, the solution was filtered by using a grade-4 crucible followed by washing with water. The crude product was purified by column chromatography by using neutral alumina with a 1:1 ratio of acetonitrile and toluene. The orange-red band was collected and analyzed. The final product was a fine dark orange solid. Yield: 0.08g (62.5%); Found: C, 53.2; H, 2.67; N, 13.2 (calculated for $C_{56}F_{12}H_{34}N_{12}OP_2Ru$: C, 52.4; H, 3.0; N, 13.1); MS, m/z: 1136.9 [M⁺- PF₆] and 993.2 [M⁺- 2PF₆],; ¹H NMR (CD₃CN, δ ppm): 9.65 (d, 6H, J=6.87 Hz), 9.05 (d, 4H, J=8.02 Hz), 8.43 (m, 6H), 8.23 (dd, 4H, J=3.44 Hz), 8.15 (d, 2H, J=9.16 Hz), 7.82 (m, 4H), 7.69 (dd, 2H, J=5.73 Hz), 7.20 (d, 2H, J=9.16 Hz) and 3.88 (s, 3H); IR (ν , cm⁻¹):3386.03 (secondary N-H), 2954.87 (sp^3 C-H), 1458.91 (C=C), 1354.91 (C=N), 1022.27 (C-N) and 811.44 (Ar para-subs.)

UV-Visible titration

The interaction of ruthenium(II) complex, $[Ru(dppz)_2(p\text{-MOPIP})]^{2+}$ with CT-DNA was analyzed using UV-Vis spectroscopy. The UV-Vis titration of the complex in a buffer was performed at room temperature using a fixed concentration of the complex with increasing concentration of DNA. The concentration of the $[Ru(dppz)_2(p\text{-MOPIP})]^{2+}$ solution was 15 μ M while the concentration of DNA titrated varied from 0 to 86 μ M. The complex-DNA solution was allowed to incubate for 10 minutes before the spectra were recorded. The titration processes were repeated several times until no more changes were observed in the spectra which indicated that the binding saturation was achieved. The changes in the concentration of the ruthenium(II) complex due to dilution at the end of each titration were negligible.

Emission titration

The intrinsic fluorescence emission spectra of the complex in Tris-buffer was carried out by using a fixed ruthenium(II) concentration to which increments of the DNA stock solution were added. Ruthenium-DNA solution was allowed to incubate for 5 minutes before the emission spectra were recorded. The excitation wavelength was set at 460 nm.

Viscosity measurements

Viscosity measurements were carried out with an Cannon Flask viscometer, suspended vertically and immersed in a water bath with a constant temperature of 25 °C. The flow time was measured with a digital stopwatch and each sample was tested three times to obtain the average flow time. Data were represented as $(\eta/\eta_0)^{1/3}$ versus binding ratio, 1/R where η is the viscosity of DNA in the presence of the complex. Specific viscosity values were calculated by the equation $\eta = (t - t_0)/t_0$, where t and t_0 represent flow time of the DNA solution and the blank buffer, respectively.

Results and Discussion

The reaction of $[Ru(dppz)_2Cl_2]$ with p-MOPIP in a ratio 1:1 treated with potassium fluorophosphate resulted in formation of $[Ru(dppz)_2p$ -MOPIP]²⁺ in PF₆ salt (Scheme 1). The purified complex was obtained as an orange-red fine solid with 62.5% yield. Conversion to chloride salts was done with the treatment of the complex in acetonitrile solution with tetra-N-butylammoniumchloride. The complex in PF₆ salt was easily soluble in acetonitrile while as a chloride salt, it was soluble in water.

Scheme 1. Preparation of [Ru(dppz)₂p-MOPIP]²⁺

Electrospray ionization mass spectrometry (ESI-MS) in acetonitrile showed a peak at m/z 1136.9 for [M⁺] while the intense peak at m/z 496.1 served as the base peak for [M²⁺]. The ¹H NMR spectrum of [Ru(dppz)₂p-MOPIP]²⁺ (Figure 1) in deuterated acetonitrile were properly defined and integrated. There are nine signals which corresponded to the protons of dppz and p-MOPIP ligands in the complex. The resonance at 9.65 ppm corresponded to the proton attached to the carbon adjacent to a coordinated nitrogen atom in the aromatic ring while the proton attached to the neighboring oxygen atom was shielded at 7.20 ppm. The proton signal for the methyl group was observed at 3.88 ppm as a singlet peak. Unfortunately, the signal that corresponded to the proton directly attached to nitrogen atom in p-MOPIP was not observed in the spectrum, probably due to fast exchange chemistry of the active proton in the imidazole ring or it was obscured by the nitrogen quadrapole broadening property [12]. However, the presence of NH was confirmed by the IR spectrum.

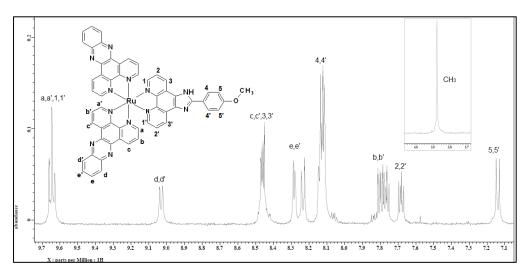


Figure 1. ¹H NMR spectrum of [Ru(dppz)₂p-MOPIP]²⁺

The IR spectrums of the complex were recorded in the region of $4000 - 280 \text{ cm}^{-1}$ and the data is summarized in Table 1. A broad and strong peak which gives information on secondary amine group of imidazole $\nu(N-H)$ was

observed at 3386 cm⁻¹. The carbon-nitrogen bonds absorptions v(C=N) and v(C-N) were at 1638 and 1354 cm⁻¹, respectively. The stretching frequency of v(C-O) at 1022 cm⁻¹ were indicative of the O-CH₃ group. In addition, a strong out of plane v(C-H) bending frequency appeared at 811 cm⁻¹ indicating that the substitution in the aromatic phenyl referred to the *para*-position. The accumulated MS, NMR and IR data supported by elemental analysis was consistent with the proposed structure and composition of the complex.

Wave number (cm ⁻¹)	Intensity
3386	strong, broad
2954	strong, sharp
1638	medium, sharp
1458	strong, sharp
1354	strong, sharp
1022	medium
811	medium
	3386 2954 1638 1458 1354 1022

Table 1. IR data of [Ru(dppz)₂p-MOPIP]²⁺

The UV-Visible absorption spectrum for $[Ru(dppz)_2p\text{-MOPIP}]^{2+}$ was recorded in acetonitrile, with a concentration of $10 \,\mu\text{M}$ at room temperature. Through comparison to related molecules [8, 9], the bands centered at around 281 nm ($\varepsilon = 66600 \, \text{M}^{-1} \, \text{cm}^{-1}$) and 358 nm ($\varepsilon = 17400 \, \text{M}^{-1} \, \text{cm}^{-1}$) were assigned to intra-ligand $\pi \to \pi^*$ transitions in the scaffold. The lowest energy band at 460 nm ($\varepsilon = 10200 \, \text{M}^{-1} \, \text{cm}^{-1}$) was attributed to the metal-to-ligand charge transfer (MLCT) transition of the $Ru(d_{\pi}) \to ligand$. The luminescence spectra of the complex upon MLCT excitation was performed at room temperature and the excitation wavelength was set at 460 nm. Figure 2 displays the emission spectra of complex in acetone (**blue**) and water (**red**) at room temperature to show the effect of the protic solvent on the complex. The complex displayed high luminescence in acetonitrile but almost no emission in water probably due to the presence of the dppz ligands in the complex. In an aqueous environment, the free electron on the nitrogen of the phenazine ring can form hydrogen bonds with water molecules [13,14]. As a result, the excited-state luminescence was quenched [15, 16]. This property is advantageous in DNA-binding studies using an aqueous buffer, where the complex binds to DNA, and the dppz ligands were protected from water, thus intense luminescence could be observed. This phenomenon has been termed as the 'molecular light switch' effect [17].

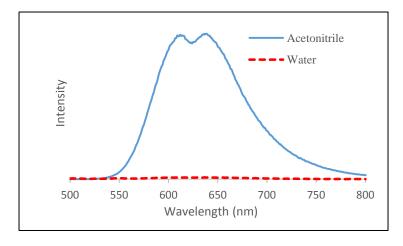


Figure 2. Luminescence spectra of [Ru(dppz)₂p-MOPIP]²⁺ in acetonitrile and water

DNA-binding studies

The interaction of the ruthenium(II) complex, $[Ru(dppz)_2(p-MOPIP)]^{2+}$ with CT-DNA was analyzed using UV-Vis spectroscopy. The UV-Vis titration of the complex in a buffer was performed at room temperature using a fixed concentration of the complex with increasing concentration of DNA.

The concentration of the $[Ru(dppz)_2(p\text{-MOPIP})]^{2+}$ solution was at 15 μ M while the concentration of DNA varried from 0 to 86 μ M. The complex-DNA solution was allowed to incubate for 10 minutes before the spectra were recorded in order to allow the interaction between DNA and the complex. The titration processes were repeated several times until no more changes were observed in the spectra indicating that the binding saturation was achieved. The changes in the ruthenium(II) complex concentration due to dilution at the end of each titration were too small and thus were negligible [18]. The absorption spectrum of complex with increasing concentration of CT-DNA is shown in Figure 3.

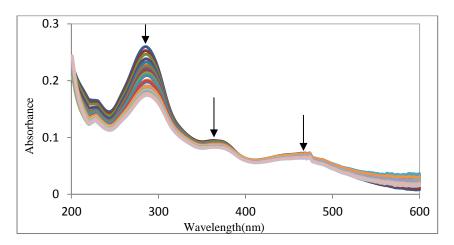


Figure 3. Absorption spectra of ruthenium(II) complex in Tris-HCl buffer upon addition of CT-DNA. [Complex] = 15uM, [DNA] = 0-86 uM

The arrows showed the absorbance changing upon the increase in DNA concentration. Normally, the hypochromic and bathochromic effects are seen during the binding process which is due to the perturbation of the metal centred MLCT band and the ligand centred π - π * band of the complex [19]. For this case, although there was no obvious red shift, hypochromicities was observed. The hypochromicities in the absorption bands for $[Ru(dppz)_2(p\text{-MOPIP})]^{2+}$ complex by the addition of CT-DNA in aqueous buffer was 29 %. This hypochromism effect has been attributed to the presence of synergic non-covalent interactions: electrostatic, groove or intercalation binding along the outside of DNA helix, and often the extent of hypochromism was parallel to the intercalative binding strength [20].

There were three well resolved bands at 284 nm, 363 nm and 474 nm. The absorption bands decreased with higher concentration of CT-DNA which fit a characteristic feature of CT-DNA binding interaction. The ultraviolet bands at 284 nm and 363 nm correlated to the intra-ligand π - π * transition [19]. The lowest energy bands at 474 nm was assigned to the MLCT transitions pf Ru(d $_{\pi}$) \rightarrow ligand according to the other related ruthenium (II) complexes [21].

In order to quantitatively illustrate the DNA binding strength of the complex, the intrinsic binding constant, K_b of complex with CT-DNA was determined by monitoring the changes of absorbance at 284 nm while increasing the concentration of DNA according to the following equation 1 [22]:

$$[DNA]/(\varepsilon_a - \varepsilon_f) = [DNA]/(\varepsilon_b - \varepsilon_f) + 1/K_b(\varepsilon_b - \varepsilon_f)$$
(1)

where [DNA] is the concentration of CT-DNA in base pairs, the apparent absorption coefficient, ϵ_a , ϵ_f and ϵ_b correspond to $A_{observed}$ /[Ru], the extinction coefficient for the free ruthenium complex and the extinction coefficient for the free ruthenium complex in the fully bound form, respectively .

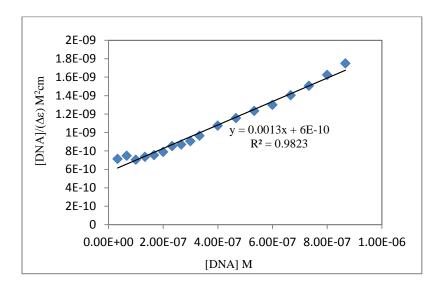


Figure 4. [DNA]/($\Delta\epsilon$) vs [DNA] for the titration of DNA with the ruthenium(II)complex

 K_b was determined by the ratio of slope to the intercept and the value obtained was 1.67 x 10^6 M⁻¹. By comparing with other similar ruthenium(II) complexes that contained p-MOPIP as ligand, this value was greater compared to $[Ru(bpy)_2(p\text{-MOPIP})^{2+}$ and $[Ru(phen)_2(p\text{-MOPIP})]^{2+}$ where the value of K_b were 5.0 x 10^5 M⁻¹ and 0.20 x 10^5 M⁻¹ [10, 13], respectively. This signified that $[Ru(dppz)_2(p\text{-MOPIP})]^{2+}$ bound more tightly to DNA compared to those complexes. In this case, the ancillary ligand dppz might have played an important influence suggesting that the high DNA binding affinity was relative to the dimension of π -conjugated aromatic area of ancillary ligands. This was mainly induced by the extended-aromatic structure of the ancillary ligand which was dppz that increased the action between the complexes and DNA. The dppz itself might also react with DNA consequently giving a higher value of binding constant as mentioned before.

The K_b value was consistent with previous report by Sun et al. [23] on a closely related compound consisting of two dppz ligands in one system, $[Ru(dppz)_2(L^1)]^{2+}$, $(L^1=5,5^2-di(1-(trimethylammonio)methyl)-2,2^2-dipyridyl cation) showed the effect of the intercalating ligand in DNA binding, with <math>K_b$ values at 0.58 x 10^6 M⁻¹. The distinct structure of *p*-MOPIP which comprised of an extended aromatic area assisted $[Ru(dppz)_2(p-MOPIP)]^2$ to bind strongly to DNA through intercalation compared to $[Ru(dppz)_2(L^1)]^{2+}$ where L^1 contained longer alkyl groups in the ligand. The binding affinity of $[Ru(dppz)_2(p-MOPIP)]^2$ was improved by bis-intercalation due to presence of more than one intercalating ligand in one system, namely *p*-MOPIP and two sites of dppz [8,14].

The intrinsic fluorescence emission spectra of complex in Tris buffer was carried out by using a fixed ruthenium(II) concentration to which increments of the DNA stock solution were added. Ruthenium-DNA solution was allowed to incubate for 10 minutes before the emission spectra were recorded. The excitation wavelength was set at 460 nm.

In the emission spectra (Figure 5), the arrow showed the emission intensity changes upon increasing the DNA concentration. The concentrations of the ruthenium(II) complex and CT-DNA were 30 μ M and 0-200 μ M, respectively. The interaction of the complex with DNA shows increased emission intensity in buffer solution. However, the emission changes were not so pronounced even after excess addition of DNA because of solvent quenching and enhancement of emission intensity when bound intercalatively to DNA. This implied that the

ruthenium(II) complex could strongly interact with DNA and be protected by DNA efficiently. In other words, the hydrophobic environment inside the DNA helix reduced the accessibility of solvent water molecules to the complex and the complex mobility was restricted at the binding site [24], leading to a decrease of the vibrational modes of relaxation. Increase in emission was an indication of intercalation between CT-DNA and the complex.

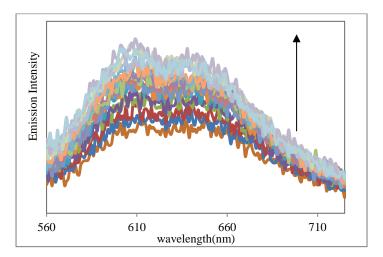


Figure 5. The emission spectra of the ruthenium(II) complex in buffer in the presence of increasing amounts of CT-DNA, $[Ru] = 30\mu M$, $[DNA] = 0 - 200\mu M$

This result also revealed the selective nature of $[Ru(dppz)_2(p-MOPIP)]^{2+}$ as a molecular switch, responding sensitively to the suitable changes in the structure of the helix, a phenomenon called the 'molecular light switch' effect, due to presence of the dppz ligands.

In order to elucidate the binding mode of [Ru(dppz)₂(*p*-MOPIP)]²⁺, viscosity measurements were carried out by keeping a constant concentration of CT-DNA and varying the concentration of the complex. Viscosity measurements provided reliable evidence to study the mode of interaction. Electrostatic or groove binding generally had no significant consequence on the viscosity of DNA, whereas classical intercalation affects the lengthening of DNA helix as base pairs are separated to accommodate foreign molecules, therefore, increasing the viscosity of DNA [25]. As shown in Figure 6, with continuing addition of [Ru(dppz)₂(*p*-MOPIP)]²⁺ to CT-DNA solution, the viscosity of CT-DNA increased. This result validated the hypothesis that [Ru(dppz)₂(*p*-MOPIP)]²⁺ bound to DNA *via* intercalation. This result was in agreement with other previous reports on ruthenium(II) polypyridyl complexes [10, 26].

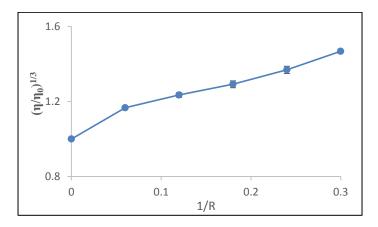


Figure 6. Effect of increasing concentration of [Ru(dppz)₂(p-MOPIP)]²⁺ on the viscosity of CT-DNA

Conclusion

In this study, a novel complex $[Ru(dppz)_2(p-MOPIP)]^{2+}$ has been synthesized. The structure and composition was supported by analysis $via^{-1}H-NMR$, mass spectrometry, elemental analysis, FT-IR, and UV-Vis spectroscopy. The binding properties of CT-DNA with the complex were studied by UV-Vis titration and viscosity measurements showed that the interaction occurred via intercalation mode giving the intrinsic binding constant of $K_b = 1.67 \times 10^6 M^{-1}$. $[Ru(dppz)_2(p-MOPIP)]^{2+}$ was bound more tightly than similar related complexes. From the comparison, it was clear that the extended aromaticity of the ancillary ligand influenced the binding strength of ruthenium(II) complex to DNA. In the luminescence titration, it was found that the $[Ru(dppz)_2(p-MOPIP)]^{2+}$ was a highly sensitive spectroscopic reporter of double-helical DNA. Whereas in aqueous solution, the luminescence was detectable only when $[Ru(dppz)_2(p-MOPIP)]^{2+}$ intercalated in the nucleic acid structure and was perhaps shielded by DNA. Therefore, $[Ru(dppz)_2(p-MOPIP)]^{2+}$ was concluded that served as a true 'molecular light switch' for the DNA structures. The stark contrast between the 'off' and 'on' states of $[Ru(dppz)_2(p-MOPIP)]^{2+}$ made this complex useful for the quantitative study of DNA and other non-polar microenvironments, a significant outcome for the design of novel intercalator reagents. However, the details of binding of complex with the sequence of DNA is not clear yet and further studies are required.

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