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- 2 Double-chain Surfactant Co(III) Complexes Containing Imidazo[4,5-
- **3 f]**[1,10]**phenanthroline and Dipyrido**[3,2-d:2'-3'-f]**quinoxaline ligands**:
- 4 Experimental and Theoretical Study
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8 9 Karuppiah Nagaraj^a, Gunasekaran Velmurugan^b, Subramanian Sakthinathan^a, Ponnambalam Venuvanalingam^b and Sankaralingam Arunachalam^a*

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A new class of surfactant Co(III) complexes, cis-[Co(ip)₂(C₁₂H₂₅NH₂)₂](ClO₄)₃ (1) and cis-10 $[Co(dpq)_2(C_{12}H_{25}NH_2)_2](ClO_4)_3$ (2) (ip = imidazo[4,5-f][1,10]phenanthroline, dpg = 11 dipyrido[3,2-d:2'-3'-f]quinoxaline), have been synthesized and characterized by various 12 spectroscopic and physico-chemical techniques. The critical micelle concentration (CMC) values 13 14 of these complexes in aqueous solution were obtained from conductance measurements. The specific conductivity data (at 303, 308, 313, 318 and 323K) served for the evaluation of the 15 temperature-dependent CMC and the thermodynamics of micellization (ΔG^0_m , ΔH^0_m and ΔS^0_m). 16 The trend in DNA-binding affinities and the spectral properties of a series of complexes, cis-17 $[Co(ip)_2(C_{12}H_{25}NH_2)_2](ClO_4)_3$ (1) and cis- $[Co(dpq)_2(C_{12}H_{25}NH_2)_2](ClO_4)_3$ (2), have been 18 experimentally and theoretically investigated. The experimental results indicate that the size and 19 20 shape of the intercalated ligand and hydrophobicity of the complexes have a marked effect on the binding affinity of complexes to CT DNA in intercalation mode, and the order of their intrinsic 21 DNA-binding constants K_b is $K_b(1) < K_b(2)$. In addition, the influence of the extended aromatic 22 ring and optical properties of the complexes can be reasonably explained by applying the DFT 23 calculations. The energy gap between HOMO and LUMO indicates that these complexes are 24 prone to interact with CT DNA. Further, molecular docking calculations have also been 25 performed to understand the nature of binding of the complexes and the result confirms that the 26 complexes interact with CT DNA through the alkyl chain. The cytotoxic activity of these 27 complexes on human liver carcinoma cancer cells were determined adopting MTT assay and 28 specific staining techniques, which revealed that the viability of the cells, thus treated was 29

significantly decreased and the cells succumbed to apoptosis as seen in the changes in the
 nuclear morphology and cytoplasmic features.

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36 Introduction

37 The understanding of the physical nature of DNA cationic surfactant interaction has primary importance due to its significance in biomedical applications, particularly as efficient gene 38 delivery vectors [1]. In cationic detergents between DNA into discrete particles each consisting 39 of a single nucleic acid molecule [2,3], which is released fast into cells with accompanying 40 decondensation. This interesting feature of DNA-surfactant interaction has prompted biophysical 41 characterization of complexes of large DNA molecules with various surfactants [4,5]. Many 42 factors are reported to affect efficient gene transfection including the type of nanoscale structure 43 and the surface charge of DNA/surfactant complex. It is realized that strong DNA/cationic 44 surfactant interaction can help to compact DNA, yielding complexes of small size [6-9]. The 45 cooperative binding of cationic surfactants on DNA chains is primarily driven by both 46 electrostatic attraction and hydrophobic effect. The cationic surfactant can be varied by the 47 molecular structures including conventional single-tailed surfactants [10], Gemini surfactants 48 49 [11], double-tailed surfactants [12], etc.

50 The mode of action of many drugs in clinical use for the treatment of cancer, genetic disorders, and viral diseases is believed to be based on their binding to nucleic acids and 51 subsequent modification of the genetic material [13]. Ribonucleic acid (RNA) also plays 52 53 essential role in normal biological process as well as in the progression of many diseases that led to growing interests in exploiting in designing of a new class of therapeutics [14-16]. A number 54 of small organic molecules are endowed with cytotoxic properties that are exerted through highly 55 specific but non-covalent and reversible interactions with nucleic acid structures [17,18]. The 56 57 non-covalent interactions are further classified in terms of intercalation and groove binding. Intercalation occurs when planar aromatic molecules are sandwiched between two adjacent base 58

59 pairs of nucleic acids. Non-intercalative binding occurs in the major or minor grooves of nucleic acids without inserting any part of the binding molecule between base pairs. The physical and 60 molecular basis of binding of natural alkaloids to nucleic acids has been a subject of extensive 61 study in the recent past [19-21]. The intercalating ligands constitute one of the most widely 62 studied groups as they form important class of compounds for cancer therapy [22]. We have been 63 interested in the synthesis and micelle forming properties of Co(III)/cu(II) complexes containing 64 lipophilic ligands [23-33]. As in biology, such compounds may exhibit novel physical and 65 chemical properties with interesting and useful associated applications. In metal-based drugs, the 66 metal can coordinate ligands in a precise three-dimensional configuration, thus allowing the 67 tailoring of the molecule to recognize and interact with a specific molecular target [34-36]. This 68 is further enhanced by different chemical modifications of ligands. Moreover, metal complexes 69 easily undergo redox reactions and ligand substitution which allow them to participate in 70 biological redox chemistry and interact with biological molecules. It is remarkable that 71 investigations in this area are focused on the use of biologically active complexes formed by 72 essential ions, such as Co (III). In spite of the great effort and success in the study of surfactant 73 Co(III) complexes, such complexes still attract much attention due to the relative simplicity of 74 their synthesis and their interesting properties. Co(III) complexes of bipyridine and 75 phenanthroline chelators are of great interest since they exhibit numerous biological properties 76 such as antitumor, anticandida and antibacterial activity [37-39]. In order to guide the design and 77 78 synthesis of new complexes with excellent bioactivity, some important factors affecting DNA binding affinities of such a kind of polypyridyl complexes, e.g., the planarity and substituted 79 phenanthroline properties of the intercalative ligands, as well as ancillary ligand effects, have 80 been experimentally summarized in certain scale [40,41a]. In addition to the experimental 81 82 studies, the transition metal polypyridyl complexes have also attracted many theoretical chemists. Various theoretical researchers have been trying to correlate some theoretical 83 predictions to the experimental findings [41b and 41c]. 84

In this surfactant Co(III) complexes, cis-85 paper, we report two $[Co(ip)_2(C_{12}H_{25}NH_2)_2](ClO_4)_3$ (1) and cis- $[Co(dpq)_2(C_{12}H_{25}NH_2)_2](ClO_4)_3$ (2) are synthesized 86 and characterized (Scheme 1 and 2). The trend in DNA-binding affinities and the spectral 87 properties of a series of surfactant Co(III) polypyridyl complexes were experimentally and 88 theoretically studied. The DNA binding constants K_b of the complexes were determined 89

systematically with spectrophotometric titration. DFT calculations were performed to rationally explain the binding trend and the molecular docking studies carried out to obtain detailed binding information of the complexes with CT DNA. This paper is mainly focused on experimentally revealing the trend in DNA binding affinities of such a type of surfactant Co(III) polypyridyl complexes and on theoretically attempting to understand them in order to effectively control the DNA binding affinities of the complexes by selecting some suitable modified phenanthrolne ligands. This structural feature offers the chance for them to be a candidate for DNA-binding

97 reagents. The cytotoxicty of these surfactant Co(III) complexes against human liver carcinoma
98 cancer cells are also reported.

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100 Results and discussion

101 Critical micelle concentration (CMC) and thermodynamics of micellization

102 The specific conductivity of the surfactant Co(III) complexes increases with the complex 103 concentration and temperature. When plots are made of [complex] versus specific conductivity, the slope is reduced after a particular value of concentration. This particular value of 104 105 concentration at which slope of the plot changes shows micellization and this concentration is chosen as CMC. Fig. 1 and SI (Supplementary Information) Fig. 1 illustrates that the plot for the 106 complexes (1) and (2). It was observed that the CMC values increased with increase in the 107 temperature for a given system. This behavior may be related to two competitive effects. First, an 108 increase in temperature causes a decrease in hydration in the hydrophilic group, which favors 109 micellization. Second, an increase in temperature also disrupts the water surrounding the 110 hydrophobic group, and this retards micellization. The relative magnitude of these two opposing 111 effects will determine CMC behavior. The study of CMC versus temperature is often undertaken 112 to obtain information on hydrophobic and head group interactions. This involves deriving 113 various thermodynamic parameters of micelle formation [56-61]. The thermodynamic 114 parameters of micellization for the surfactant Co(III) complexes are presented in Tables 1 and SI 115 Table 1. The observed more negative Gibbs free energy of micellization indicates more favored 116 micellization for the system under study. Moreover, since the changes of CMC with temperature 117 are small, the value of ΔH^0_m and ΔS^0_m must be rather inaccurate and should be considered as 118

only approximate. As mentioned in our previous reports [23,25,28,29,31], the CMC values for surfactant Co(III) complexes in the present study were also very low compared to those of the simple organic surfactants. Thus it is suggested that our metal surfactant complexes have more capacity to associate themselves forming aggregates than the ordinary synthetic organic surfactants. Moreover, introduction of a metal complex to the hydrophilic part of the amphiphile can remarkably enhance the ability of aggregation.

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126 **DNA binding studies**

127 Electronic absorption spectral studies

Titration with UV spectroscopy is an effective method to examine the binding mode of nucleic 128 acid with metal complexes since the observed changes of the spectra may give evidence of the 129 existing interaction mode [62]. Thus, in order to provide evidence for the possibility of binding 130 131 of complexes to DNA, spectroscopic titration of the solutions of the surfactant Co(III) complexes with DNA has been performed. The UV spectra have been recorded for a constant DNA 132 concentration in different [compound/DNA] mixing ratios (r) and are shown in Fig. 2 and 3 and 133 SI Fig. 3 and 4. As seen from the Figures. 2,3 and SI Figures 3,4 with the increase of the 134 135 concentration of CT DNA, the absorption spectrum of surfactant Co(III) complex (1) shows hypochromism 28% for below CMC and 31% for above CMC and complex (2) shows 136 137 hypochromism 34% for below CMC and 37% for above CMC with slight red shift on the addition of increasing amounts of DNA. Hypochromism is suggested to arise due to an 138 139 intercalative mode of binding involving a strong stacking interaction between extending aromaticity of ligand and the base pairs of DNA [63]. Also Co(III) complex containing long 140 aliphatic chain enhances hydrophobic interaction strongly with intercalating the base pairs of 141 DNA. The observed spectroscopic changes are thus, consistent with intercalation of complexes 142 143 into the DNA base stacks. In order to further elucidate quantitatively the affinity of the surfactant Co(III) complex to CT-DNA, the intrinsic binding constant, K_b has been determined using the 144 equation [64], 145

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$$[DNA]/(\varepsilon_a - \varepsilon_f) = [DNA]/(\varepsilon_0 - \varepsilon_f) + 1/K_b(\varepsilon_0 - \varepsilon_f)$$

147 where, [DNA] is the concentration of DNA expressed in base pairs ; ε_a , ε_f and ε_0 are the 148 apparent, free and fully bound Co(III) complex extinction coefficients. A plot of [DNA]/ (ε_a - ε_f) 149 versus [DNA] gives K_b as the ratio of the slope to intercept.

The K_b values of the surfactant Co(III) complexes with CT DNA are given in Table 2. In 150 151 complex (1), which contains ip ligand, that is expected to be planar and possesses a smaller π system than dpg, intercalation is lower [65-68]. Since the complex (2), contains dpg ligand (2.4 \times 152 10^{6} M⁻¹ for below CMC and 3.1×10^{6} M⁻¹ for above CMC), it would provide an aromatic moiety 153 extending from the metal center through which overlapping would occur with the base pairs of 154 DNA by intercalation. Besides the binding constant at below CMC values are lower than that at 155 above the CMC values. Above cmc value the surfactant complex presents as both monomers as 156 157 well as micelles. Aggregation of complex molecules into micelles would reduce their ability to bind to DNA. So there is a difference in K_b values at above cmc and below cmc values. Also the 158 K_b of surfactant Co(III) complexes is very much higher than that for the ordinary metal 159 complexes, like $[Co(bpy)_3]^{3+}$ (K_b, 9.3 ×10³ M⁻¹) [69], $[Co(bpy)_2(imp)]^{3+}$ (K_b, 1.1 ×10⁴ M⁻¹) [70], 160 $[Co(bpy)_3(BHBMe)]^{2+}$ (K_b, 1.23 ×10⁴ M⁻¹), $[Co(bpy)_2(BHBNO_2)]^{2+}$ (K_b, 2.06 ×10⁴ M⁻¹) [71], 161 $[Co(ip)_2Cl_2]Cl (K_b, 2.9 \times 10^3 \text{ M}^{-1}), [Co(dpq)_2Cl_2]Cl (K_b, 6.4 \times 10^3 \text{ M}^{-1}) [72].$ This implies that this 162 series of the complexes can intercalate between the base pairs of DNA and thus be protected by 163 DNA efficiently, since the hydrophobic environment inside the DNA helix reduces the 164 accessibility of solvent water molecules to the complex and the complex mobility is restricted at 165 166 the binding site. Such a trend is also in agreement with that in the binding constants (K_b).

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168 Trend in DNA-binding affinities of the complexes and theoretical explanations

169 As reported in the literature DNA molecule is generally an electron-donor and the intercalated complex is an electron-acceptor and there are π - π stacking interactions in the DNA-binding of 170 171 these complexes in intercalation mode. Higher HOMO energy of DNA molecule and lower LUMO energy of the complex result in a stronger interaction between DNA and the complex. 172 173 Kurita and Kobayashi [73-75] have reported a simple calculation model and reported DFT computed results for stacked DNA base-pairs with backbones. It should be a rather reasonable 174 approximation model for DNA, and thus should be useful for such a discussion. The optimized 175 geometries of the complexes (1) and (2) are given in SI Fig. 4 (SI-Supporting Information). To 176

177 gain further insight into the influence of the extended aromatic ring and trend in DNA-binding activities of the complexes the combined molecular orbital energy level graph for complexes (1) 178 179 and (2) is given (SI Fig 5). The HOMOs and LUMOs are often used to relate the spectral properties of complexes and provide decisive clues in designing new complexes. Therefore it is 180 181 essential to identify and understand the nature of various segments of the complex and their individual contributions towards HOMOs and LUMOs. Hence, the contribution of various 182 183 fragments of the complexes has been analyzed using QMForge Program [76]. The complexes have been segmented into three fragments, namely Co (Cobalt), L1 (equatorial ligand), L2 (alkyl 184 chain) and their corresponding percentage contributions to frontier molecular orbitals are 185 summarized in Table 3. 186

The complex (2) has the lowest band gap (3.07 eV) due to the extension of aromatic ring 187 188 of the L1 destabilizing the HOMO levels. The frontier molecular orbital shows that the HOMO of complex (1) and (2) is mainly localized on the L2 (93% and 99% respectively). The LUMO of 189 190 complex (1), there is mainly centered on the L1 (85%) whereas in complex (2) it is mainly localized on Co (dvz 54%) and L1 (43%). On the other hand, the HOMO-1 to HOMO-3 of 191 complex (2) is localized on L2 of the complex. LUMO+1 of the complex (1) is mainly 192 composed of 53% Co (d_{vz}) and 45% of L1. However, extension of aromatic ring in the axial 193 194 position have a direct effect on the orbital energy of HOMO and as the result HOMO of complex (2) is more stabilized than that of complex (1). The LUMOs are not influenced by this ligand. 195

The calculated results indicate that the extension of aromatic ring in the axial position can destabilize the energy of HOMO more significantly than those of the LUMO resulting in narrower HOMO-LUMO energy gap. The LUMO of the complexes of (1) and (2) lie at -6.92eV and -6.86eV respectively, which distribute predominantly on the intercalative ligands. The predominant LUMO population on the intercalative ligands of the complexes should be advantageous to accept the electron from base-pairs of DNA and the trend in the K_b is well correlated with the LUMO energy values of the complexes.

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204 Molecular docking details

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205 The above experimental results show that these surfactant Co(III) complexes bind to DNA in the intercalation mode and the DNA-binding affinities of these complexes increase in the order: (1) 206 207 < (2). Obviously, there are subtle but detectable differences occurred in the DNA-binding properties. Such differences may be resulted from the changes of geometric and electronic 208 209 structures of these complexes (especially the intercalative ligand) due to the different modified phenanthrolne on the intercalative ligand. Although the single crystal structures of these 210 211 surfactant Co(III) complexes have not been obtained as far, the DFT calculations can give us some important parameters of the geometric structures of these complexes, as well as some 212

useful information on the electronic structures.

To understand the efficiency of a biologically active drug molecule, the knowledge of its 214 215 binding location in the CT DNA is very essential and significant. Moreover, the docking study further corroborated the existence of interactions between the Co(III) complexes and DNA. Thus 216 the molecular docking calculations were performed and the most probable docked poses are 217 given in Fig. 4. As seen from the Fig. 4 clearly shows that both the complexes could fit well into 218 219 the CT DNA with a binding site of three base pairs, preferentially involving the G-C residues as 220 revealed by the docked structure. The availability of the extended alkyl chain makes the complex 221 could bind well to the hydrophobic interior of CT DNA. The complex (2) interacts with CT DNA through intercalation and gains additional stabilization due to aromatic moiety extending 222 223 from the metal center. Thus, the complex (2) is found to have higher binding affinity than complex (1). In both the cases, the surfactant Co(III) complexes holding three base pairs is like 224 225 A-T, G-C, G-C to the outer surface, i.e., the groove of the CT DNA. Thus, the G-C regions seem to facilitate a better fit of the complexes into the CT DNA. It is interesting to note that the 226 binding energy of complex (1) was 12.0 kcal mol⁻¹ less than that of complex (2) due to the π - π 227 stacking interaction of extending of aromatic moiety. The aromatic ring structure of surfactant 228 Co(III) complex having a long alkyl chain allows for torsional rotation in order to fit into the 229 narrower helical curvature of the CT DNA. This excellently agrees with the outcome of other 230 231 experiments (binding constant and spectroscopic studies). Thus, our molecular modeling studies throw light on the binding modes by which these complexes interact with CT DNA and 232 complement the experimental observations. 233

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235 Thermal denaturation studies

The dissociation of a duplex nucleic acid into two single strands results in significant 236 hyperchromism around 260 nm. The binding of a metal complex to a nucleic acid induces a 237 conformational change of the latter to alter so-called denaturation temperatures depending on the 238 strength and mode of interactions between complex and nucleic acid. In the present work the 239 melting curves of CT-DNA in the absence and presence of complex (1) and (2) are presented in 240 SI Fig. 6. A T_m of CT-DNA in buffer was determined as for surfactant complexes under our 241 experimental conditions (74.0 \pm 0.2[°] C). The DNA intrinsic binding constant of the surfactant 242 Co(III) complexes at T_m were calculated by using McGhee's equation (Eq. (4))[77], where T_m^0 is 243 the melting temperature of CT-DNA alone, T_m is the melting temperature in the presence of the 244 Co(III) complexes, ΔH_m is the enthalpy of DNA (per base pair), R is the gas constant, K is the 245 DNA-binding constant at T_m, L is the free complex concentration (approximate by the total 246 complex concentration) at T_m, and n is the size of the binding site. 247

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$$1/T_{m}^{0} - 1/T_{m} = (R/\Delta H_{m}) [ln(1+KL)]^{1/n}$$
 (4)

By substitution of the required parameters into Eq. (1), K was determined to be 6.68 x 249 10^5 M^{-1} for the title complex at 85° C. Only few thermodynamic parameters such as free energy, 250 enthalpy and entropy changes upon binding of metal complexes to DNA have been measured, 251 although there have been many reports on the interaction of metal complexes with DNA. In fact, 252 253 the thermodynamic parameter of DNA-complex formation is essential for a thorough understanding of driving forces of the binding of metal complexes to DNA [78]. The change of 254 standard enthalpy was determined according to the van't Hoff's equation (Eq. (5)). The changes 255 of standard free energy and standard entropy of the binding of the title complex to DNA were 256 determined according to Eqs. (6) and (7), where K₁ and K₂ are the DNA-binding constants of the 257 complex at the temperatures T₁ and T₂, respectively. ΔG^0_T , ΔH^0 , and ΔS^0 are the changes of 258 standard free energy, standard enthalpy, and standard entropy of the binding of the titled 259 complexes to CT-DNA, respectively. 260

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$$\ln(K_1/K_2) = H^0/R (T_1 - T_2/T_1T_2)$$
 (5)

$$\Delta G^0_T = -RT \ln K$$

(6)

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(7)

$$\Delta G^0_{T} = \Delta H^0 - T \Delta S^0$$

The value of ΔH^0 is derived to be -82.8 k J mol⁻¹ by substituting K₁ = 1.17 x 10⁶ M⁻¹ (T1 264 = 298K) and $K_2 = 6.68 \times 10^5 \text{ M}^{-1}$ (T₂ = 358K) into Eq. (4). By substituting $K_1 = 1.17 \times 10^6 \text{ M}^{-1}$ 265 1 (T1 = 298K) and ΔH^{0} = -82.8 k J mol⁻¹ into Eqs. (5) and (6), $\Delta G^{0}_{298 \text{ K}}$ = -34.6 k J mol⁻¹ and ΔS^{0} 266 = -16.2 J mol⁻¹ K⁻¹ at 25[°] C were derived. It is clearly observed from the experimental results that 267 the complex formation for all the cases was spontaneous with similar negative ΔG^0 values. The 268 negative binding free energy suggests that the energy of the complex-DNA adduct is lower than 269 the sum of the energies of the free complex and DNA, and the binding of the titled complexes to 270 CT DNA is favorable at room temperature. The possible explanation of the entropically driven 271 DNA binding of these types of metal complexes has been discussed in details from the viewpoint 272 273 of molecular interaction by Hag *et al.* [79] and by us earlier. The negative entropy implies that the degree of freedom of the titled complexes and DNA conformation is reduced upon complex-274 275 DNA binding. According to the thermodynamic data, interpreted as follows, the model of 276 interaction between a drug and biomolecule can be concluded as [80]: (1) $\Delta H < 0$ and $\Delta S < 0$, 277 hydrophobic forces; (2) Δ H>0 and Δ S>0, van der Waals interactions and hydrogen bonds; (3) Δ H>0 and Δ S<0, electrostatic interactions [81]. In order to elucidate the interaction of our 278 279 complex with DNA, the thermodynamic parameters were calculated. When we apply this analysis to the binding of the complex with CT DNA, we find that $\Delta H < 0$ and $\Delta S < 0$. 280 Therefore, intercalations via hydrophobic interactions are probably the main forces in the 281 binding of the investigated complexes to CT DNA. These results provide that these surfactant 282 Co(III) complexes may interacts with CT DNA in an intercalation mode. 283

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285 Competitive binding between ethidium bromide (EB) and surfactant Co(III) complexes

In order to investigate the mode of binding of the surfactant Co(III) complexes to DNA, the competitive binding experiment using ethidium bromide $(2 \times 10^{-5} \text{ M})$ was carried out. Ethidium bromide (EB) is one of the most sensitive fluorescent probes that can bind to DNA [82-83]. The emission spectra of EB bound to DNA in the absence and the presence of the surfactant Co(III) complexes (1) and (2) are given in Fig. 5 and SI Fig. 8. The addition of these complexes to DNA pretreated with EB caused an appreciable reduction in the emission intensity, indicating that the

replacement of the EB fluorophore by the complex results in a decrease of the binding constantof ethidium bromide to DNA. According to the classical Stern-Volmer equation [84]:

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$$I_0/I = 1 + K_{sv}[Q]$$

where I₀ and I are the fluorescence intensities in the absence and the presence of the complex, 295 respectively. K_{sv} is a linear Stern–Volmer constant and r is the ratio of the total concentration of 296 the complex to that of DNA. The fluorescence quenching of EB bound to DNA by the surfactant 297 Co(III) complexes (1) and (2) are shown in SI Fig. 8. The quenching plots illustrate that the 298 299 quenching of EB bound to DNA by the surfactant Co(III) complexes are in good agreement with the linear Stern-Volmer equation, which also indicates that the complexes bind to DNA. In the 300 plot of Io/I versus [complex]/[DNA], K_{sv} is given by the ratio of the slope to intercept. The K_{sv} 301 values for our surfactant Co(III) complexes (1) and (2) are 0.157 and 0.183, respectively. These 302 data suggest that the interaction of surfactant Co(III) complex (2) with DNA is stronger than 303 complex (1), which is consistent with the spectral results described above. 304

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306 Electrochemical studies

307 The electrochemical investigations of metal-DNA interactions can provide a useful complement to spectroscopic methods, e.g., for non-absorbing species, and yield information about 308 interactions with both the reduced and oxidized form of the metal [85]. In general, the 309 electrochemical potential of a small molecule will shift positively when it intercalates into DNA 310 double helix, and it will shift in a negative direction in the case of electrostatic interaction with 311 DNA [86]. The complete scan in the range +1.0 V to -1.5 V of surfactant Co(III) complexes in 312 the absence of DNA shows a cathodic wave at -0.72 V and an anodic wave at 0.01 V for 313 complex (1) and a cathodic wave at -0.24 V and an anodic wave at 0.06 V for complex (2) (SI 314 Figures 9 and 10). This quasi-reversible wave can be assigned to the couple Co(III)/Co(II). 315 Complexes (1) and (2) have similar behavior and the corresponding potentials are given in Table 316 5. The quasi-reversible redox couple Co(II)/Co(III) for each complex in buffer solution has been 317 studied upon addition of CT DNA and the corresponding potentials as well as their positive 318 shifts are given in Table 4. No new redox peaks appeared after the addition of CT DNA to each 319 320 complex, but the current intensity of all the peaks decreased significantly, suggesting the

existence of an interaction between each complex and CT DNA. The decrease of current intensity can be attributed to an equilibrium mixture of free and DNA bound complex to the electrode surface [86]. For increasing amounts of CT DNA, the cathodic potential E_{pc} for all complexes shows a positive shift. Hence the positive shifts in the CV peak potential of surfactant

325 Co(III) complexes are indicative of intercalation of these complexes into the DNA [86].

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327 Circular Dichroism Studies

CD spectra is a useful technique in diagnosing changes in DNA morphology during drug–DNA 328 interactions as the CD signals are quite sensitive to the mode of DNA interactions with small 329 molecules [87]. The characteristic CD spectra of right-handed B form DNA consist of two 330 bands: a positive band (275 nm) due to the base stacking and a negative band (245 nm) due to 331 the right-handed helicity. The observed changes of CD signals of DNA are usually assigned to 332 the corresponding changes in DNA structure [87]. It is generally accepted that the classical 333 intercalation enhances the base stacking and stabilizes helicity, and thus increases intensities of 334 the both bands, whereas simple groove binding and electrostatic interaction of small molecules 335 show less or no perturbation on the base stacking and helicity bands [88]. In SI Fig. 11, the CD 336 spectrum of CT DNA was monitored in the presence of increasing amounts of complexes (1) and 337 (2). The positive band showed an increase in molar ellipticity with a red shift of the band 338 maxima when the complex concentration was progressively increased. This increase in intensity 339 340 with a red shift in positive bands suggests that surfactant Co(III) complexes binds to CT DNA via intercalation. 341

342

343 Viscosity measurements

The binding modes of the surfactant Co(III) complexes were further investigated by viscosity measurements. In the absence of crystallographic structure data, hydrodynamic methods, which are sensitive to increase in DNA length, are regarded as the least ambiguous and the most critical tests of binding in solution [88a]. Optical or photophysical probes generally provide necessary, but not sufficient clues to support an intercalative binding model. Under appropriate conditions 349 intercalation causes a significant increase in the viscosity of DNA solutions due to the separation of base pairs at intercalation sites and, hence, increase the overall DNA contour length whereas 350 351 ligands that bind exclusively in the DNA grooves, groove-face or electrostatic interactions typically cause a bend (or kink) in DNA helix reducing its effective length and thereby its 352 viscosity. The effects of the surfactant Co(III) complexes (1) and (2) on the viscosity of DNA 353 are shown in SI Fig. 12. Upon addition of increasing the amounts of the titled complexes, the 354 355 relative viscosity of DNA increase steadily. These results suggest that the titled surfactant Co(III) complexes intercalates between the base pairs of DNA, in agreement with the above 356 experimental results. 357

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359 Cytotoxity Studies

360 MTT assay

We examined the cytotoxicity of the effects of these surfactant Co(III) complexes on cultured HepG2 liver cancer cells by exposing cells for 24 and 48 h to the medium containing the complex1 at $8.3-74.7\mu m$ (10-90µg/mL) and complex 2 at $8.0-72.0\mu m$ (10-90µg/mL) concentration. *In vitro* antitumor activity of these compounds was determined according to the percentage of nonviable cells (%NVC) which was calculated by the following equation:

366 NVC% = [number of NVC/total number of cells] x 100

The results of these experiments are summarized in Table 5. As shown in table increasing the concentration of surfactant Co(III) complexes was accompanied by progressive increase in the NVC %. This is due to the fact that by increasing the concentration of cationic surfactant complexes the adsorption of ions on cell membranes increases, leading to increase in penetration and antitumor activity.

The results of the cytotoxic activity on human tumor cell lines was determined according to the dose values of drug exposure required to reduce survival in the cell lines to 50% (IC₅₀). It should be noted that the action of the complex as antitumor agents is found to be dependent on the type of tumor cell line (HepG2) tested but, as shown from the results, surfactant Co(III) complexes show excellent cytotoxic activity against tumor cell lines (HepG2) and, at very low concentrations, reduces the survival to 50%. This is due to the fact that cobalt complexes have a

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378 capacity to reduce the energy status in tumors as well as to enhance tumor hypoxia [88b and 88c], which also influences their antitumor activities. It may be also concluded that the level of 379 380 cellular damage inflicted by these complex depends on the nature of their axial ligands. It is known that phenanthroline-containing metal complexes have a wide range of biological activities 381 382 such as antitumor, antifungal, apoptosis [89,90] and interaction with DNA inhibiting replication, transcription and other nuclear functions and arresting cancer cell proliferation so as to arrest 383 384 tumor growth. In general, the high selectivity of action by redox-active cobalt complexes upon tumors is due to their specific reactivity [91]. From these results, surfactant Co(III) complexes 385 seems to offer promise due to the high electron affinity of the metal (which increases its ability 386 to bind DNA) and the ready reducibility of the compounds [92]. 387

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Fluorescence Microscopic Analysis of apoptotic cell death (AO and EB staining)

390 AO/EB staining adopting fluorescence microscopy also revealed apoptosis from the perspective of fluorescence. After HepG2 liver cancer cells were exposed to the concentrations of surfactant 391 392 Co(III) complexes for 24 h. In this study, we used acridine orange/ethidium bromide (AO/EB) double staining assay [93]. Acridine orange is taken up by both viable and nonviable cells and 393 emits green fluorescence if interrelated into double stranded nucleic acid (DNA). Ethidium 394 bromide is taken up only by nonviable cells and emits red fluorescence by intercalation into 395 DNA. We distinguished four types of cells according to the fluorescence emission and the 396 morphological aspect of chromatin condensation in the stained nuclei: (1) viable cells showing 397 light green fluorescing nuclei with highly organized structure; (2) early apoptotic cells having 398 bright green fluorescing nuclei with chromatin condensation and nuclear fragments; (3) late 399 apoptotic cells having orange to red fluorescing nuclei with condensed or fragmented chromatin; 400 401 and (4) necrotic cells having red fluorescing without chromatin fragmentation. Viable cells have uniform bright green nuclei with organized structure. Apoptotic cells have orange to red nuclei 402 with condensed or fragmented chromatin. Necrotic cells have a uniformly orange to red nuclei 403 with condensed structure (Fig. 6). Our results indicate that surfactant Co(III) complexes induced 404 apoptosis at the concentrations evaluated, in agreement with the cytotoxic results. The results 405 suggest that the complex treatment caused more cells to take to death in HepG2 liver cancer 406 407 cells.

408

409 Apoptosis Detection Hoechst 33342 DNA Staining

It is possible to perform apoptosis detection assay with Hoechst 33342 (Sigma B-2262), but the 410 411 increase in fluorescence seen in the apoptotic cells may be less dramatic. Hoechst dyes can also be obtained from Molecular Probes. H33342 is a "vital" DNA stain that binds preferentially to 412 413 A-T base-pairs. The cells require no permeabilization for labeling, but do require physiologic conditions since the dye internalization is an active transport process. This condition typically 414 415 varies among cell types (Stander et al., 2009). To investigate if HepG2 liver cancer cells were triggered to undergo apoptosis due to the exposure of surfactant Co(III) complexes, 416 417 morphological changes of apoptosis were performed in the treated cells by Hoechst 33342 staining. Apoptosis is one of the major pathways that lead to the process of cell death. After the 418 cells were treated with IC₅₀ concentrations of surfactant Co(III) complexes for 24 and 48 h the 419 cells were observed for cytological changes adopting Hoechst 33342 staining. The observations 420 421 revealed that the complexes brought about cytological changes such as chromatin fragmentation, binucleation, cytoplasmic vacuolation, nuclear swelling, cytoplasmic blabbing and late apoptosis 422 indication of dot-like chromatin and condensation (Fig. 7) whereas untreated cells did not show 423 such changes. Data collected from the manual counting of cells with normal and abnormal 424 425 nuclear features. Both apoptotic and necrotic cells increased in dose-dependent manner. These data clearly indicated that higher doses of surfactant Co(III) complexes resulted in remarkable 426 chromatin condensation and nuclear fragmentation in HepG2 liver cancer cells. 427

428

429 Conclusions

In this work the surfactant Co(III) complexes containing polypyridyl ligand has been synthesized and successfully characterized. The cmc values of these surfactant Co(III) complexes are very low showing more capacity to associate themselves, forming aggregates, compared to those of ordinary synthetic organic surfactants. The experimental results indicate that these surfactant Co(III) complexes could binds to CT DNA by intercalative mode and the binding constant of complex (1) is smaller than that of complex (2). Computations support this and further it reveals that the intercalative binding between surfactant Co(III) complexes and CT DNA is due to the presence of extending aromaticity of ligand and a long aliphatic chain of the complexes. Furthermore, it was observed that surfactant Co(III) complex exhibits considerable ability to bind with the G-C-rich sequence of DNA. As the aromatic moiety extending from the metal centre, the binding ability increases. Both the surfactant Co(III) complexes are cytotoxic to human liver carcinoma cancer cells. Therefore, these complexes might prove to be of application in target-based cancer therapy since the mechanism of cell death appears to be essentially apoptosis, but necrosis also is one of the desired endpoints in cancer therapy.

444 Experimental Section

445 Materials and methods

All the reagents were of analytical grade (Sigma-Aldrich and Merck). Calf thymus DNA 446 447 obtained from Sigma-Aldrich, Germany, was used as such. The spectroscopic titration was carried out in the buffer (50 mM NaCl-5 mM Tris-HCl, pH 7.1) at room temperature. A solution 448 of calf thymus DNA in the buffer gave a ratio of UV absorbance ~1.8-1.9:1 at 260 and 280 nm, 449 indicating that the DNA was sufficiently free of protein [38]. Milli-Q water was used to prepare 450 451 the solutions. The ligands, ip and dpq were synthesized by using previously published procedures [39,40]. The precursor Co(III) complexes, cis-[Co(LL)₂Cl₂]Cl (where LL₂ = ip and 452 453 dpq) were prepared by a similar method to the previously published reports [41,42]. Absorption spectra were recorded on a UV-Vis Spectrophotometer using cuvettes of 1 cm path length, and 454 455 emission spectra were recorded on a JASCO FP 770 spectrofluorimeter. FT-IR spectra were recorded on a FT-IR Perkin Elmer spectrophotometer with samples prepared as KBr pellets. ¹H 456 457 NMR spectra were recorded on a BRUKER 500 MHz Spectrometer using DMSO as solvent. 458 Conductivity studies were carried out on aqueous solutions of the complexes with an Elico 459 conductivity bridge type CM 82 and a dip-type cell with a cell constant of 1.0. The carbon, 460 hydrogen and nitrogen content of the complexes were determined on a CHNS Elemental Analyzer-elementar (Model: Vario EL III) available at Sophisticated Analytical Instrument 461 Facilities (SAIF), Central Electrochemical Research Institute (CECRI) Karaikudi, India. The 462 anticancer studies were carried out at PCBS Research Centre, Pondicherry University. 463

464 Synthesis of double chain surfactant Co(III) complexes

465 Preparation of cis- $[Co(ip)_2(C_{12}H_{25}NH_2)_2](ClO_4)_3$ (1)

To a solution of cis-[Co(ip)₂Cl₂]Cl (0.53 g) dissolved in water (10 mL), dodecylamine 466 (0.26 mL) in ethanol (2 mL) was added dropwise over a period of 30 min. The mixture was set 467 468 aside at room temperature for 4 days until no further change was observed. Afterwards, a saturated solution of sodium perchlorate in perchloric acid was added. A pasty solid mass 469 470 separated out slowly and it was filtered off, washed with small amounts of alcohol followed by acetone, and then dried in air. The semi-dried material was further dried over fused calcium 471 472 chloride in a drying pistol and stored in a vacuum desiccator. UV λ_{max} (CH₂Cl₂)/nm 248 (π - π *), 290 (n- π^*), 370 (MLCT). FT-IR (KBr)/ cm⁻¹ 786, 725, 1384 (C=N), 2914, 2852, 1104 and 621. 473 ¹H NMR (400 MHz; CDCl₃): δ 0.96 (m, 6H, CH₃), 1.29-4.02 (44H, Aliphatic-H), 4.21 (m, 2H, 474 NH₂), 4.41(m, 2H, NH₂), 5.43 (m, 2H, NH), 7.34 (m, 12H, Ar-H), 7.67 (s, 2H, Ar-H). ¹³C NMR 475 (400 MHz; CDCl₃): δ 24.27, 24.30, 29.32, 32.92, 33.12, 34.77, 35.01 (Aliph-C's), 117.83, 476 118.58, 118.81, 120.92, 127.37, 127.86, 128.35, 129.35, 129.79, 137.17, 138.27, 148.08, 150.08, 477 160.30, 161.60, 165.37, 165.67 (Ar-C's). Calcd. for C₅₃H₇₈CoN₁₀: C, 52.60, H, 6.07, N 11.75. 478 Found: C 51.60, H, 6.30, N, 11.83. 479

480 Preparation of cis- $[Co(dpq)_2(C_{12}H_{25}NH_2)_2](ClO_4)_3$ (2)

The cis- $[Co(dpq)_2(C_{12}H_{25}NH_2)_2](ClO_4)_3$ complex was synthesized similar to the 481 482 procedure described above, except that we have added dpg ligand in the reaction mixture instead of ip ligand. [Caution: Perchlorate salts of metal complexes with organic ligands are potentially 483 484 explosive. Only small amounts of the material should be prepared and handled with great care]. UV λ_{max} (CH₂Cl₂)/nm 250 (π - π^*), 293 (n- π^*), 374 (MLCT). FT-IR (KBr)/ cm⁻¹ 791, 728, 1386 485 (C=N), 2912, 2850, 1100 and 620. ¹H NMR (400 MHz; CDCl₃): δ 0.98 (m, 6H, CH₃), 1.18-4.31 486 (44H, Aliphatic-H), 3.72-3.82 (m, 4H, NH₂), 5.43 (m, 2H, NH), 6.5-7.8 (16H, Ar-H). ¹³C NMR 487 488 (400 MHz; CDCl₃): δ 24.27, 24.30, 29.32, 32.92, 33.12, 34.77, 35.01 (Aliph-C's), 117.83, 118.58, 118.81, 120.92, 127.37, 127.86, 128.35, 129.35, 129.79, 137.17, 138.27, 148.08, 150.08, 489 490 160.30, 161.60, 165.37, 165.67 (Ar-C's). Calcd. for C₅₅H₇₈CoN₁₀: C, 52.62, H, 6.11, N 11.38. Found: C 52.67, H, 6.17, N, 11.59. 491

492 Estimation of amount of Cobalt in Complexes 1 and 2

493 Cobalt content in the surfactant Co(III) complexes was estimated according to Kitson 494 [47]. Briefly, a known weight of the complex was reduced with tin and concentrated 495 hydrochloric acid. The reduced aqueous cobalt(II) ion was made up to 10 mL in volumetric flask using 0.1 M perchloric acid. 2 mL of this solution and 1 mL of 50% ammonium thiocyanate
solution were pipetted out into a 10 mL volumetric flask and made up to the mark with acetone.
The absorbance of this solution was measured at 625 nm against a reagent blank. From the
absorbance, the concentration of cobalt was calculated. The percentage of cobalt thus obtained
for our surfactant Co(III) complexes is Co 4.98 (4.89) for complex (1) and Co 4.88 (4.71) for
complex (2).

502 Estimation of critical micelle concentration (CMC) values

The CMC values of the complexes were determined conductometrically using a specific 503 conductivity meter. The conductivity cell was calibrated with KCl solution in the appropriate 504 concentration range. Various concentrations of surfactant Co(III) complexes were prepared in the 505 506 appropriate range in aqueous solution. The conductivity of these solutions was measured at 303, 308, 313, 318 and 323K. The temperature of the thermostat was maintained constant, to be 507 within ± 0.01 . The conductance was measured after thorough mixing and temperature 508 equilibration at each dilution. The establishment of equilibrium was checked by taking a series of 509 readings at 15 min intervals until no significant change occurred. At all temperatures a break in 510 the conductance versus concentration in the plots, characteristic of micelle formation, was 511 observed. The CMC values were determined by fitting the data points above and below the break 512 to two equations of the form y = mx + c and solving the two equations simultaneously to obtain 513 the point of intersection. Least-squares analysis was employed, and the correlation coefficients 514 were greater than 0.98 in all the cases. 515

516 **DNA binding experiments**

The DNA concentration per nucleotide was determined adopting absorption spectroscopy 517 using the known molar extinction coefficient value of 6600 M^{-1} cm⁻¹ at 260 nm [48]. Absorption 518 519 titrations were performed by using a fixed surfactant Co(III) complex concentration to which increments of the DNA stock solution were added. Surfactant Co(III) complex-DNA solutions 520 521 were incubated for 10 min before the absorption spectra were recorded. For fluorescence experiments, DNA was pretreated with ethidium bromide (EB) for 30 min. The surfactant 522 523 Co(III) complexes were then added to this mixture and the effect on the emission intensity was measured. The samples were excited at 515 nm and emission was observed between 500 and 700 524 nm. These experiments were carried out in 50 mM NaCl-5 mM Tris-HCl at pH 7.1 in aqueous 525 media. All voltammetric experiments were performed in a single compartment cell with a three-526

527 electrode configuration on a EG&G PAR 273 potentiostat equipped with a personal computer. The working electrode was a glassy carbon and the reference electrode was standard calomel 528 529 electrode. A platinum wire was used as the counter electrode. The supporting electrolyte was phosphate buffer at pH 7.1. Solutions were deoxygenated by purging with nitrogen gas for 15 530 531 min prior to measurements; during measurements a stream of nitrogen gas was passed over the solution. Circular dichroic experiments were recorded on a JASCO J-716 spectropolarimeter 532 533 (220-320 nm) were obtained at 25°C using a quartz cell of 1cm path length. Each CD spectrum was collected after averaging over at least 4 accumulations using a scan speed of 100 nm min⁻¹ 534 and 1s response time. The region between 220nm and 320 nm was scanned for each sample. The 535 spectra was recorded in the absence and in the presence of surfactant Co(III) complexes. 536 537 Viscosity experiments were carried out using an Ubbelodhe viscometer maintained at a constant temperature of $29.0 \pm 0.1^{\circ}$ C. Calf thymus DNA (1×10⁻⁵M) sample solutions were prepared by 538 sonication in order to minimize complexities arising from DNA flexibility. Data were presented 539 as $(n / n^0)^{1/3}$ versus binding ratio, where g is the viscosity of DNA in the presence of the 540 complex, and η^0 is the viscosity of DNA alone. The relative viscosity was calculated according 541 to the relation $\eta = (t - t_0)/t_0$, where t_0 is the flow time for the buffer and t is the observed flow 542 543 time for DNA in the presence and absence of the complex [49].

544

545 Cytotoxicity assay

The cytotoxicity of the surfactant Co(III) complexes were measured in the MTT (3-(4.5-546 547 dimethylthiazol-2-yl)-2,5-diphenyl tetrazolium bromide) assay as described by us earlier [50]. The complexes were first dissolved quantitatively in dimethyl sulfoxide (DMSO, Sigma, USA) 548 to make the stock solution. Briefly, cells were seeded at a density of 5 x 10^4 HepG-2 liver cancer 549 cells/well into 96-well plates. After 24 h, the cells were treated with surfactant cobalt (III) 550 551 complexes at various concentration (10, 30, 60, 90 µg/ml) and incubated for 24 and 48 hours as indicated. At the end of the incubations, 10µl of 3-(4-5 dimethylthiozol-2-yl) 2-5 diphenyl-552 553 tetrazolium bromide (MTT) (5 mg/ml) per well was added and incubated in dark at 37°C for 4 hours. The formazan crystals formed after 4 hours were solubilized in 100µl of DMSO after 554 aspirating the medium. The absorbance was monitored at 570 nm (measurement) and 630 nm 555 using a 96 well plate reader (Bio-Rad, Hercules, CA, USA). The IC₅₀ value was defined as the 556 concentration of compound that produced a 50% reduction of cell viability. 557

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555

559 Evaluation of apoptosis (Acridine orange and ethidium bromide staining)

Acridine orange and ethidium bromide staining was performed as described by Spector et 560 al. [50]. Twenty-five microliters of cell suspension of each sample (both attached, released by 561 trypsinization, and floating), containing 5×10^5 cells, was treated with AO and EB solution (one 562 part of 100 mg/mL AO and one part of 100 mg/mL EO in PBS) and examined under a 563 fluorescent microscope (Carl Zeiss, Germany) using an UV filter (450-490 nm). Three hundred 564 cells per sample were counted in tetraplicates for each dose point. Cells were scored as viable, 565 apoptotic or necrotic as judged by the staining, nuclear morphology and membrane integrity, and 566 percentages of apoptotic and necrotic cells were then calculated. Morphological changes were 567 568 also observed and photographed. The amount of 200 μ L of dye mixture (100 μ L/mg AO and 100 µL/mg EB in distilled water) was mixed with 2 mL cell suspension (30,000 cells/mL) in 6-well 569 plate. The suspension was immediately examined and viewed under Olympus inverted 570 fluorescence microscope (Ti-Eclipse). We observed untreated cells as controls and cells treated 571 572 with testing material IC₅₀ concentrations for 24 h of exposure. HepG2 were seeded in a 24-well plate (50,000 cells per well). After 24 h of cells incubation, the medium was replaced with 100 573 574 μ L medium containing IC₅₀ dose of testing material. Untreated cells served as the control. After 24 h, aspirate the media and treat with prepared dye and observe under the fluorescent 575 576 microscope.

577

578 Hoechst 33342 (H342) staining

Stock H342 solution should be originally suspended in dH₂O at a 1mM concentration 579 (H342 will precipitate in PBS). This procedure is very sensitive to cell concentration and pH of 580 the media. Cells should be approximately 1-2x10 /ml, in buffered media, pH 7.2. It is also 581 helpful to include 2% fetal calf serum to maintain the cells. Add H342 dye from the stock 582 solution to cell suspension to respective final concentration. Cells are then incubated at 37 C for 583 1 hour. Drug was added and incubated for 24 and 48 hours. Time is a critical factor due to the 584 transport of the dye. Typically, 30 minutes is a minimum, but it is important to remember that the 585 signal may begin to degrade after ~120 minutes. It is recommended that the staining kinetics be 586

empirically defined. Analyze apoptosis under fluorescent microscope after incubation. Washingis not recommended.

589

590 **Computational details**

Computational studies on the complexes have been performed using HEX 6.3 software 591 which is an interactive molecular graphics program for the interaction, docking calculations, and 592 to identify possible binding site of the biomolecules [51]. The density functional theory (DFT) 593 calculations at B3LYP level were carried out using the GAUSSIAN09 program [52]. The 594 geometries of the complexes under study were optimized using standard 6-31G** basis sets for 595 N, C and H elements and LANL2DZ for Cobalt with effective core pseudo potentials [53]. The 596 coordinates of metal complexes were taken from their optimized structure as a .mol file and were 597 598 converted to pdb format using GaussView software. The crystal structure of B-DNA (PDB ID: 1BNA) is retrieved from the protein data bank [54]. All possible poses have been considered as 599 600 starting points and the docking analysis was performed. The default parameters were used for the docking calculation with correlation type shape only, FFT mode at 3D level, grid dimension of 6 601 602 with receptor range 180 and ligand range 180 with twist range 360 and distance range. Visualization of the docked systems has been further analyzed with PyMOL software package 603 604 [55].

605

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758		Scheme captions
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761	Figur	e. 1. Electrical conductivity vs. complex (1) concentration in aqueous solutions.
762	Figur	e. 2. Absorption spectra of complex (1) (Above cmc): in the absence (dotted line) and in the
763	prese	nce (solid lines) of increasing amounts of CT DNA. {Inset: Plot of [DNA] / $(\epsilon_a - \epsilon_f)$ vs.
764	[DNA]}. [complex] = 1.0×10^{-4} M; [DNA] = $0.9.1 \times 10^{-5}$ M.

Figure. 3. Absorption spectra of complex (1) (Below cmc): in the absence (dotted line) and in the

presence (solid lines) of increasing amounts of CT DNA. {Inset: Plot of [DNA] / $(\epsilon_a - \epsilon_f)$ vs.

767 [DNA]}. [complex] = 1.0×10^{-6} M; [DNA] = $0.9.1 \times 10^{-5}$ M.

- Figure. 4. Docked pose of surfactant Co(III) Complexes bound to the CT DNA: (a) Complex (1)
- with B-DNA (b) Complex (2) with B-DNA.
- 770

Figure. 5. Emission spectra of EB bound to CT DNA: in the absence and in the presence of surfactant Co(III) complex (1). [EB] = 2×10^{-5} M, [DNA] = 1×10^{-4} M, [Complex] = $0 - 1.43 \times 10^{-6}$

Figure. 6. Photomicrographs of control and AO and EB stained HepG2 liver cancer cells incubated for 24 hours with surfactant Co(III) complexes (1) and (2). A,B Untreated control cells. C,D surfactant Co(III) complex treated control cells. ; Viable (light green), early apoptotic (bright green fluorescing), late apoptosis (red to orange fluorescing) and necrosis (red fluorescing) cells were observed. Magnification at 20x.

- Figure. 7 Surfactant Co(III) complexes (1) and (2) induces apoptosis in HepG2 liver cancer cells.
 Representative fluorescent micrographs of HepG2 liver cancer cells stained with Hoechst 33342
 fluorescent dye after the compound exposure for 24 and 48 hours. I, II Untreated control cells;
 A,B,C and, D surfactant Co(III) complex treated control cells. A, C 24 hours; B, D 48 hours.
 Magnifications at 20X.
- 783
- 784
- 785

Table captions

Table 1. CMC values of the surfactant Co(III) complex (1) in aqueous solution.

Table 2. The binding constant (K_b) and Stern-Volmer constant (K_{SV}) of surfactant Co(III) complexes (1) and (2) with CT DNA.

Table 3. Frontier molecular orbital compositions (%) of surfactant Co(III) complexes (1) and (2)
at the B3LYP/LANL2DZ level.

Table 4. Cyclic voltammetric data (mV) of surfactant Co(III) complexes (1) and (2), a scan rate
of 100 mV/s with phosphate buffer as supporting electrolyte.

Schemes

Table 5. IC₅₀ by MTT-3-(4-5 dimethylthiozol-2-yl) 2-5 diphenyl-tetrazolium bromide method.















(b)

(a)

500



Wavelength, nm

821 822

818

819



(1)

(2)



20 x

20 x

829

830

831

 Control
 24 hrs
 48 hrs

 (1)
 (A)
 (C)

 (1)
 (B)
 (C)

 (2)
 (11)
 (B)
 (D)

126

20x



833

834

835



Fig. 7

836

837

838					
839					
840					
841			Tables		
842			Table 1		
843	Temp.	CMC× 10 ⁵	$-\Delta G^{0}_{mic}$ (kJ mol ⁻¹)	$-\Delta H^{0}_{mic}(kJ mol^{-1})$	$T\Delta S^{0}_{mic}$ (kJ mol ⁻¹)
844				Lift me(no mor)	rido mie (no mor)
845	303K	9.45	35.0 ± 0.2	11.3 ± 0.3	23.8 ± 0.4
846	308K	9.64	57.3 ± 0.1	18.8 ± 0.1	38.5 ± 0.1
847	313K	9.97	78.5 ± 0.1	26.3 ± 0.2	52.2 ± 0.1
848	318K	10.5	99.9 ± 0.4	34.3 ± 0.1	65.6 ± 0.2
849					
850	323K	10.89	114.9 ± 0.4	40.3 ± 0.1	74.7 ± 0.2
851					

Table 2

Surfactant Co(III) Complexes	$K_b(M^{-1})$ Below CMC Above cmc		Hypochromism	K _{sv} (M ⁻¹))
Complex (1)	(5.7±0.2)×10 ⁵	(1.3±0.3)×10 ⁶	29	31	0.1573
Complex (2)	(2.4±0.1)×10 ⁶	(3.1±0.2)×10 ⁶	34	37	0.1829

853

852

Table 3

Energy		Complex (1)		. (Complex (2)		
Levels -	Со	L1	L2	Со	L1	L2	
НОМО-3	0.00	0.00	100.00	0.00	0.00	100.00	
НОМО-2	0.43	95.84	3.73	0.00	0.00	100.00	
HOMO-1	0.09	40.64	59.27	4.36	3.65	91.99	
НОМО	0.05	6.86	93.09	0.91	0.27	98.82	
LUMO	0.07	85.15	14.78	53.55	43.02	3.43	
LUMO+1	53.15	45.70	1.15	52.04	17.35	30.62	
LUMO+2	51.58	17.57	30.85	0.66	96.65	2.69	
LUMO+3	0.82	96.35	2.83	1.43	97.58	1.00	

Table 4	
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Surfactant Co(III) Complexes	E _{pc1} (mV)	E _{pa1} (mV)	ΔE_p (mV)	E _{1/2} (mV)	i_{pa}/i_{pc}	E _{pc2} (mV)
Complex (1)	-718.5	+13.5	+732.0	-352.5	0.6	-1208.5
Complex (1) + DNA	-714.5	+7.5	+721.9	-353.5	0.7	-1202.5
Complex (2)	-238.5	+56.5	+295.0	-91.0	1.5	-908.5
Complex (2)	-258.5	+167.5	+426.0	-45.5	1.8	-838.5
+ DNA						

Table 5

Complexes	IC ₅₀ (µ1	m)
-	24 h	48 h
(1) (2)	(6.8± 0.2) (6.0± 0.2)	(6.1± 0.2) (5.0± 0.2)

Graphical Abstract

