

New DNA binding ligands as a model of chromomycin A₃

Shuhei Imoto,^{a,b} Yoshinari Haruta,^a Kyouichi Watanabe^a and Shigeki Sasaki^{a,b,*}

^aGraduate School of Pharmaceutical Sciences, Kyushu University, 3-1-1 Maidashi, Higashi-ku, Fukuoka 812-8582, Japan

^bCREST, Japan Science and Technology Agency, Kawaguchi Center Building, 4-1-8 Honcho, Kawaguchi, Saitama 332-0012, Japan

Received 7 June 2004; revised 20 July 2004; accepted 21 July 2004

Available online 7 August 2004

Abstract—Small molecules with DNA-binding affinity within the minor groove have become of great interest. In this study, new DNA-binding ligands were designed to mimic Chromomycin A₃ (CRA₃), which contains a hydroxylated tetrahydroanthracene chromophore substituted with di and trisaccharides. The trisaccharide part of CRA₃ that is supposed to contribute to form the Mg²⁺-coordinated dimer was expected to be mimicked by a simple alkyl group attached to the chromophore part as new model compounds. The present study has successfully demonstrated that the new ligands form Mg²⁺-coordinated dimer complexes to exhibit DNA-binding affinity.

© 2004 Elsevier Ltd. All rights reserved.

Chromomycin A₃ **1** (CRA₃) is a potent antitumour antibiotic, and is a part of the aureolic acid group including plicamycin, mithramycin and olivomycins.¹ It has been shown that CRA₃ binds in the minor groove of GC-rich regions of duplex DNA to show inhibition against DNA-dependent DNA polymerases as well as RNA polymerases.² Although the aureolic family of drugs are not currently used for therapeutic purposes, the new compounds of this family are reevaluated as potential candidates in cancer therapy.³ CRA₃ contains a hydroxylated tetrahydroanthracene chromophore substituted with various sugars. Studies on the structure–activity relationship have shown that the sugar parts contribute to DNA-binding properties as well as biological activities.⁴ The aglycone of CRA₃, chromomycinone, does not bind to DNA.⁵ It has been established by structural studies with NMR and X-ray crystallography that CRA₃ binds in the minor groove of the DNA as an Mg²⁺-coordinated dimer and G-specific hydrogen bonds between CRA₃ and DNA provide the G-rich sequence specificity. It has been also shown from the crystal structure that Mg²⁺ adopts an octahedral coordination to O1 and O9 atoms of two chromophores and two water molecules, as schematically shown in **2**. The keto-phenol structure of O1 and O9 is responsible for Mg²⁺ binding, and mutual stackings of the aromatic

part of the chromophore of one monomer with the C–D glycosidic linkage of the other monomer stabilize dimer structure.⁶ Triethyleneglycol was used to mimic the C/D/E trisaccharide part for the formation of Mg²⁺-coordinated dimer, but the role of the triethyleneglycol part for DNA binding of the model compound remains unclear.⁷ We hypothesized that an alkyl chain would produce higher stacking interaction to stabilize the dimer structure complexed with Mg²⁺,⁸ and designed a new model compound **3**. In this study, the hexyl group was chosen as an alkyl chain to mimic the C–D part stacking with the aromatic part (Fig. 1). Herein we describe that the model compound **3** binds to DNA as an Mg²⁺-coordinated dimer.

The synthesis of **3** is shown in Scheme 1. We planned to construct the *peri*-hydroxy aromatic skeleton by strong-base-induced [4 + 2] cycloaddition of homophthalic anhydrides with α -sulfinyl-substituted derivatives of enolizable enones, which was developed by Kita's group.⁹ Homophthalic anhydride derivative **6** was obtained from the corresponding homophthalic acid derivative **5**.¹⁰ The α -sulfinyl-substituted cyclohexenone **8** was synthesized from commercially available (*R*)-3-methylcyclohexanone. The disulfinylated derivative **7** was obtained via a two-step reaction by regioselective sulfinylation with LDA–PhS–SPh,¹¹ followed by reaction with potassium *tert*-butoxide and PhSSO₂Ph. Oxidation of **7** with an equivalent of *m*-CPBA produced the 2-phenylthio-cyclohexenone, which was further exposed to another equivalent of *m*-CPBA to furnish the enone **8** in good

Keywords: Chromomycin A₃; DNA minor groove binder.

* Corresponding author. Tel.: +81-92-6426615; fax: +81-92-6426876; e-mail: sasaki@phar.kyushu-u.ac.jp

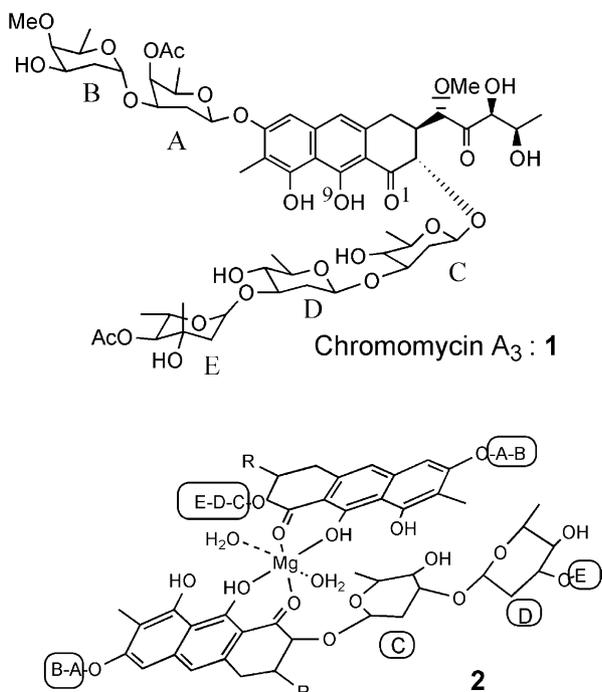


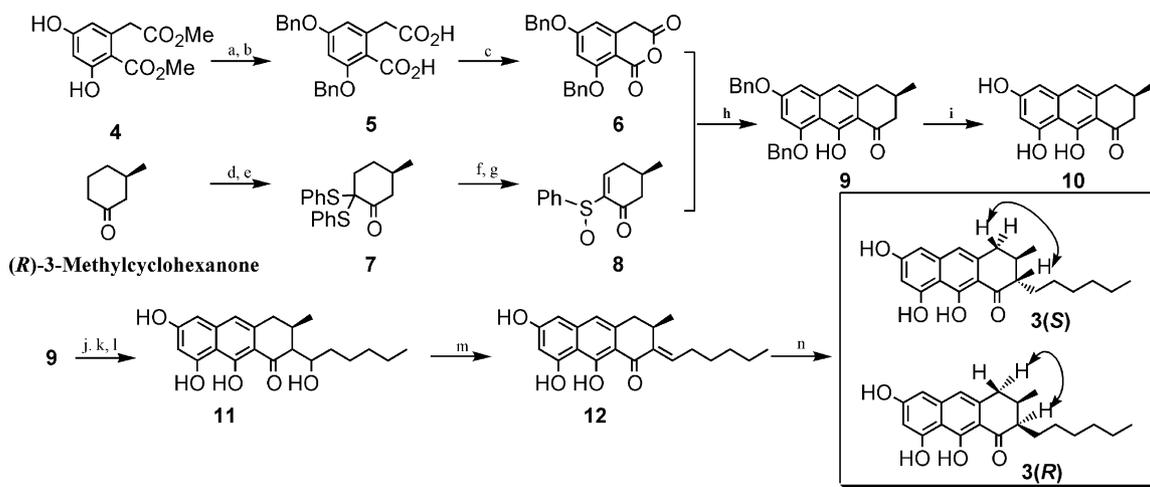
Figure 1. Chromomycin A₃ (CRA₃, **1**) and the schematic structure of Mg²⁺-coordinated dimer (**2**).

yield. The [4 + 2] cycloaddition was performed by using **6** and **8** in the presence of NaH in refluxing dioxane to produce **9** in 20–30% yield. After a number of attempts, we found that molecular sieves catalyze the [4 + 2] cycloaddition to produce **9** in good yield.¹² The hexyl group was introduced by aldol reaction, followed by deprotection of the benzyl ethers by hydrogenolysis with H₂-palladium hydroxide to give **11**. Dehydroxylation was accomplished under acidic condition, and the produced *exo*-olefin was hydrogenated with H₂-palladium hydroxide to furnish a mixture of **3(S)** and **3(R)** isomers. The diastereomers were separated by normal phase HPLC

using ChiralPak-AS as a column, and their stereochemistry was determined by ¹H COSY and NOESY spectra.¹³

The Mg²⁺-binding properties of the compounds were investigated by measurement of the UV spectra (Fig. 2) in methanol. The UV spectra of **3(S)** was changed by the addition of MgCl₂ to produce a hyperchromic effect around 420 nm (Fig. 2A) with the isosbestic points. Stoichiometry of the complex was evaluated by the Job plot, and the 2:1 **3(S)** to Mg²⁺ complex was proven (Fig. 2B). The ability of the dimer complex of **3(S)** with Mg²⁺ resembles the TEG-chromophore conjugate previously reported.⁷ It has been also shown that the **3(R)** isomer forms the 2:1 **3(R)** to Mg²⁺ complex. The UV titration data (Fig. 2A) was analyzed based on the 2:1 complex to produce a stability constant of $K_s = 1.0 \times 10^6 \text{ M}^{-1}$. In contrast, the Job plots with the nonalkylated compound **10** (Fig. 2C) were not clear compared to Figure 2B, probably because **10** forms a mixture of 1:1 and 2:1 complexes. These results have shown that the alkyl chain of **3** contributes to the formation of the dimer complex.

We also analyzed the complex compositions by positive ESI-MS measurements. The mass spectrum of **3(S)** in the presence of MgCl₂ indicated 2:1 ligand to Mg²⁺ complexes ($m/z = 707$) in addition to 1:1 complexes (429, 415, 401) (Fig. 3B). In the MS spectra of **10** with MgCl₂, there are the major peaks ($m/z = 345, 331, 317$) due to the 1:1 ligand to Mg²⁺ complexes together with the minor peak corresponding to the 2:1 complex (Fig. 3C). There is a small tendency for the formation of 2:1 complexes with Ca²⁺ (Fig. 3D). These results by the positive ESI-MS measurements agree with those obtained by UV measurements in that **3(S)** forms 2:1 complexes with MgCl₂. It is also indicated that the alkyl chain of **3(S)** plays an important role for dimer formation. The **3(R)** isomer produced similar ESI-MS spectrum.



Scheme 1. Reagents and conditions: (a) NaH, BnBr, 93%; (b) KOH, EtOH/H₂O, 96%; (c) TMS ethoxyacetylene, 91%; (d) PhS–SPh, LDA, THF, –78 °C to rt, 82%; (e) PhSSO₂Ph, *tert*-BuOK, THF, 0 °C, 77%; (f) *m*-CPBA, CH₂Cl₂, –50 °C, quant.; (g) *m*-CPBA, CH₂Cl₂, –60 °C, 73%; (h) MS 4 Å, rt, 75%; (i) Pd(OH)₂, H₂ gas, quant.; (j) TBSOTf, TEA, CH₂Cl₂, quant.; (k) hexanal, BF₃Et₂O, –70 °C, quant.; (l) Pd(OH)₂, H₂ gas, 55%; (m) TsOH, benzene, 40 °C, 56%; (n) Pd(OH)₂, H₂ gas, 96%.

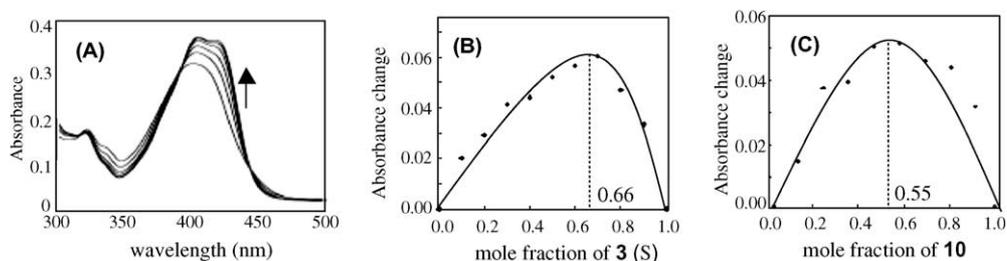


Figure 2. The UV spectrum change of 3(S) by the addition of MgCl_2 (A), and Job plot with 3(S) (B) and 10 (C).

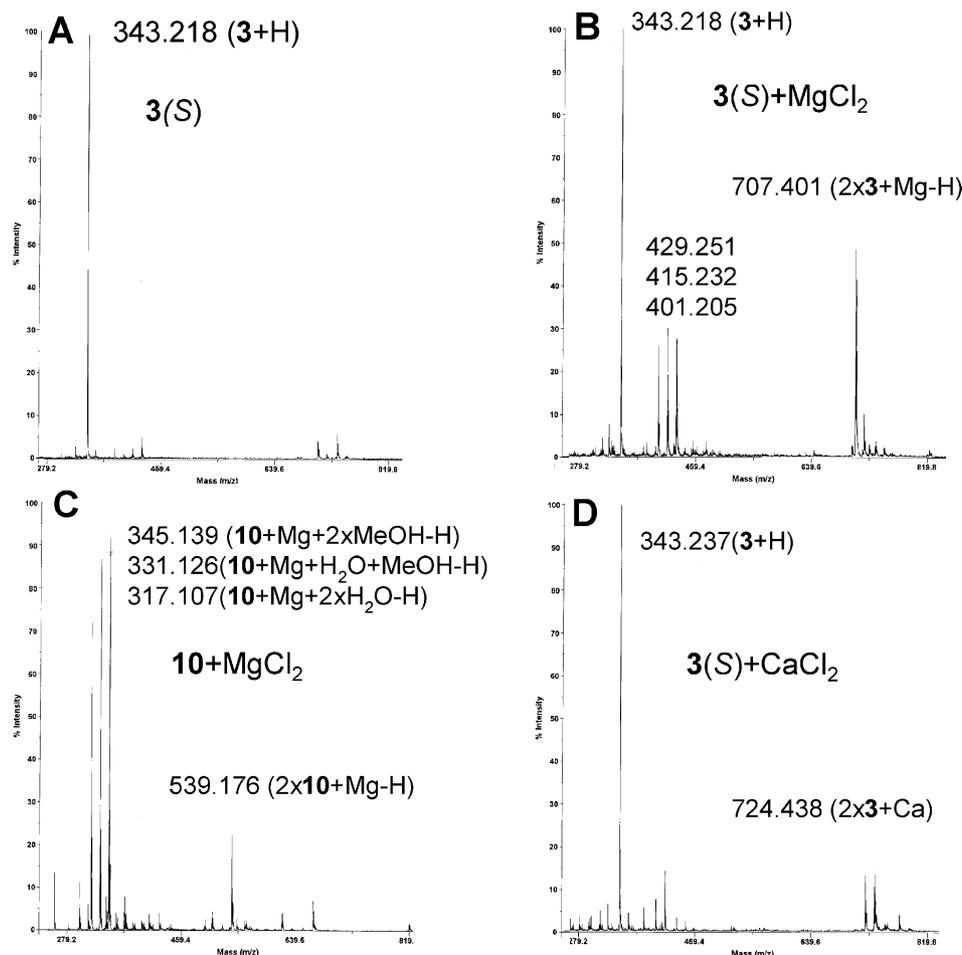


Figure 3. Analysis of the complexes by ESI-MS measurements: (A) 3(S) only; (B) 3(S) + MgCl_2 ; (C) 10 + MgCl_2 ; (D) 3(S) + CaCl_2 .

The binding affinity of the ligand with the duplex DNA was evaluated by a competitive displacement assay using fluorescent ethidium bromide (ETBr). The emission intensity of ethidium bromide is increased by binding with duplex DNA, and is inhibited by another competitive binder. Since competitive DNA binders displace DNA-bound ETBr to decrease emission regardless of their binding modes, although the binding mode of ETBr is intercalation, displacement assay with ETBr is widely used for primary estimation of the binding affinity of the compounds.¹⁴ In this study, self-complementary 12 base pair DNA duplexes, CT12 and CA12, were used for measurement of the binding affinity.¹⁵ CA12 includes an AATT region, while CT12 has a GCGC region at the middle of the duplex. Displacement assay was carried out

at 25°C using a solution of the duplex DNA (CT12 or CA12, 1 μM) in 20 mM Tris-HCl buffer. ETBr was effectively displaced by 3(S) in the buffer containing 1 mM MgCl_2 (Fig. 4A), clearly indicating the relatively high affinity of 3(S) to DNA. Conversely, displacement was not observed with use of 10 in the buffer containing MgCl_2 (Fig. 4B), or with use of 3(S) in the buffer containing NaCl instead of MgCl_2 (Fig. 4C), showing low DNA-binding affinity of the ligands under these conditions. Moderate displacements were observed with 3(S) in the buffer containing CaCl_2 (Fig. 4D). These results have indicated that the DNA-binding affinity of 3(S) correlates with the tendency of dimer complex formation mediated by Mg^{2+} , and that the alkyl chain of 3(S) is also an important contributor for DNA binding. The 3(R)

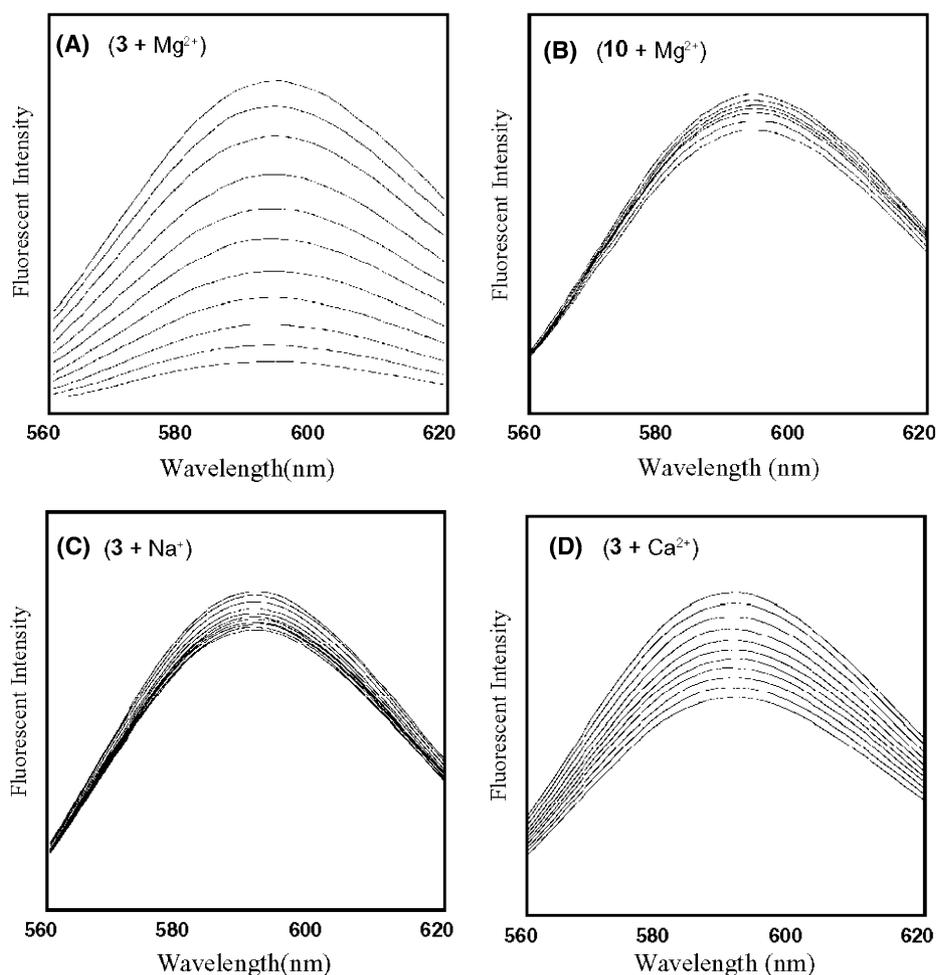


Figure 4. The ethidium bromide displacement assay with **3(S)** and **10**: (A) **3(S)** in the presence of 1 mM MgCl_2 , (B) **10** in the presence of 1 mM MgCl_2 , (C) **3(S)** in the presence of 2 mM NaCl , (D) **3(S)** in the presence of 1 mM CaCl_2 . One micromolar each of CT12 and ethidium bromide were used. The ligand concentration was increased from 0 to 10 μM . $\lambda_{\text{ex}} = 546 \text{ nm}$.

isomer showed almost the same results in the ethidium bromide displacement assay. Concentrations of the ligand to inhibit 50% of fluorescence intensity of ETBr are expressed as IC_{50} values, and correspond to the relative binding affinity of the ligand (Table 1). Selective binding of Hoechst 33258¹⁶ to the AATT region of DNA has been proven in this ethidium displacement assay (the IC_{50} values for CT12 and CA12; >20 vs 1 μM , respectively). Displacement profile of CRA_3 is somewhat complexed, because only one of two ethidium bromide molecules bound to CT12 was displaced. This is probably because the limited region of CT12 is tightly bound with CRA_3 .¹⁷ The stereochemistry of **3** did not affect the binding affinity, and the ligand **3** showed slightly weaker binding affinity to CA12 than Hoechst 33258. The IC_{50} values of the new ligands are not different towards CT12 or CA12, and sequence selectivity of the new ligands was not clear in this ethidium displacement assay.¹⁸

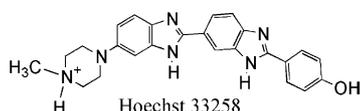


Table 1. Concentrations to displace 50% DNA-bound ethidium bromide (IC_{50} , μM)^a

Compounds	IC_{50} (μM) ^b	
	CT12 ^c	CA12 ^d
CRA_3	3 ^e	9
3(S)	3	3
3(R)	3	2.5
3(R + S)	3.5	3
10	No inhibition	No inhibition
11	6.5	4.5
12	2.5	2.5
Hoechst 33258	>20	1

^a 1 μM each of the duplex DNA and ethidium bromide were used in 20 mM Tris–HCl containing 1 mM MgCl_2 at pH 8.0. $\lambda_{\text{ex}} = 546 \text{ nm}$, $\lambda_{\text{em}} = 590 \text{ nm}$.

^b Concentration to displace 50% DNA-bound ethidium bromide is expressed as the dimer complex except for Hoechst 33258.

^c CT12: (^{5'}CGTAGCGCTACG)₂.

^d CA12: (^{5'}CGCGAATTCGCG)₂.

^e IC_{50} value to displace one of two ethidium bromide molecules bound to CT12.¹⁷

In conclusion, we have developed the new DNA binding molecules as a model of Chromomycin A₃. The model ligands **3(S)** and **3(R)** have shown a tendency to form 2:1

ligand-to-Mg²⁺ complexes in methanol. In comparison with **10** without containing the alkyl chain, it has been revealed that the alkyl chain is responsible for the formation of the dimeric complex of **3** with Mg²⁺. The DNA binding experiments have indicated that the alkyl chain of **3** plays a key role for Mg²⁺-mediated DNA binding. There is no significant difference in binding properties examined in this study between the **3(S)** and the **3(R)** isomer. The present study has shown that the simple alkyl chain can mimic some roles of the trisaccharide part for Mg²⁺-dimer complex formation and Mg²⁺-mediated DNA binding. Further efforts are currently underway to develop new ligands that bind to DNA more strongly based on the structure of **3**.

Acknowledgements

This work was supported by a Grant-in-Aid for Scientific Research (B) from Japan Society for the Promotion of Science (JSPS).

References and notes

- Cory, M. DNA Minor-Groove Binding Compounds as Antitumor Agents. In *Cancer Chemotherapeutic Agents*; Foye, W. O., Ed.; ACS Professional Reference Book; ACS: Washington, DC, 1995; Chapter 8, pp 330–331.
- Muller, W. E. G. **1977**, *1*, 457, pp 457–474.
- Remsing, L. L.; Bahadori, H. R.; Carbone, G. M.; McGuffie, E. M.; Catapano, C. V.; Rohr, J. *Biochemistry* **2003**, *42*, 8313–8324.
- Hayasaka, T.; Inoue, Y. *Biochemistry* **1969**, *8*, 2342–2347.
- Kaziro, Y.; Kamiyama, M. *J. Biochem.* **1967**, *62*, 424–429.
- Hou, M.-H.; Robinson, H.; Gao, Y.-G.; Wang, A. H.-J. *Nucl. Acids Res.* **2004**, *32*, 2214–2222.
- Silva, D. J.; Kraml, C. M.; Kahne, D. *Bioorg. Med. Chem.* **1994**, *2*, 1251–1259.
- We have reported that the alkyl chains enhance DNA-binding affinity of small molecules: Sasaki, S.; Shibata, T.; Torigoe, H.; Shibata, Y.; Maeda, M. *Nucleos. Nucleot. Nucl. Acids* **2001**, *20*, 551–558.
- Iio, K.; Ramesh, N. G.; Okajima, A.; Higuchi, K.; Fujioka, H.; Akai, S.; Kita, Y. *J. Org. Chem.* **2000**, *65*, 89–95.
- Kim, S.; Fan, G.; Lee, J.; Lee, J. J.; Kim, D. *J. Org. Chem.* **2002**, *67*, 3127–3130.
- Oppolzer, W.; Petrziilka, M. *Helv. Chim. Acta* **1978**, *61*, 2755–2762.
- Details will be published soon.
- Selected data for compound **3(S)** and **3(R)**. **3(S)**: ¹H NMR: δ (ppm) 16.64 (1H, b s), 10.02 (1H, s), 6.72 (1H, s), 6.46 (1H, d, *J* = 2.3 Hz), 6.36 (1H, d, *J* = 2.3 Hz), 3.04 (1H, dd, *J* = 16.0, 3.5 Hz), 2.59 (1H, dd, *J* = 16.0, 7.5 Hz), 2.36–2.33 (1H, m), 2.24–2.19 (1H, m), 1.89–1.83 (1H, m), 1.71–1.67 (1H, m), 1.37–1.27 (8H, m), 1.08 (3H, d, *J* = 6.7 Hz), 0.87 (3H, t, *J* = 6.9 Hz), IR (cm⁻¹, KBr) 3384, 1635, 1596, HR-ESIMS (*m/z*) calcd for C₂₁H₂₇O₄ (M+H)⁺ 343.1904, found 343.1879. **3(R)**: ¹H NMR: δ (ppm) 16.44 (1H, b s), 9.95 (1H, s), 6.74 (1H, s), 6.47 (1H, d, *J* = 2.3 Hz), 6.37 (1H, d, *J* = 2.3 Hz), 2.97 (1H, dd, *J* = 16.2, 3.6 Hz), 2.79 (1H, dd, *J* = 16.2, 6.7 Hz), 2.60 (1H, dt, *J* = 6.7, 4.0 Hz), 2.47–2.43 (1H, m), 1.88–1.84 (1H, m), 1.49–1.30 (9H, m), 0.97 (3H, d, *J* = 6.7 Hz), 0.88 (3H, t, *J* = 6.9 Hz), IR (cm⁻¹, KBr) 3384, 1635, 1596, HR-ESIMS (*m/z*) calcd for C₂₁H₂₇O₄ (M+H)⁺ 343.1904, found 343.1890. The stereochemistry of **3(S)** and **3(R)** was determined by ¹H COSY and NOESY as illustrated in Scheme 1.
- Fox, K. R., Ed.; *Drug-DNA Interaction Protocols: Optimal Absorbance and Fluorescence Techniques for Measuring DNA-Drug Interactions*; Humana Press: Totowa, NJ, 1997; Vol. 90, pp 195–218.
- Alam, M. R.; Maeda, M.; Sasaki, S. *Bioorg. Med. Chem.* **2000**, *8*, 465–473.
- A representative DNA minor groove binder with selectivity to an A₃T₃ region. Crystal structure of Hoechst 33258 bound to d(CGCAAATTTGCG)₂ duplex has been reported. Spink, N.; Brown, D. G.; Skelly, J. V.; Neidle, S. *Nucl. Acids Res.* **1994**, *22*, 1607–1612.
- Two ethidium bromide molecules bind to one CT12 or CA12 duplex. We have observed that some minor groove binders such as distamycin displace only one of two ethidium bromide molecules.¹⁵
- DNA-binding affinity of the dimer complex of **3**-Mg²⁺ was estimated from the IC₅₀ value to be ca. *K*_s = 2 × 10⁶ M⁻¹ to CT12, and 7 × 10⁶ M⁻¹ to CA12. Several attempts to estimate DNA binding by footprinting experiments were not successful so far, probably because of insufficient DNA-binding affinity of **3**. Further modification of **3** for higher affinity to DNA is now in progress.