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Ring-Closing Metathesis for the Synthesis of a Highly G-Quadruplex Selective Macrocyclic Hexaoxazole Having Enhanced Cytotoxic Potency

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Abstract

The synthesis of a 24-membered macrocyclic hexaoxazole *via* ring-closing metathesis is described. The target compound selectively stabilizes G-quadruplex DNA with no detectable stabilization of duplex DNA. An MTT cytotoxicity assay indicated that this unsaturated macrocyclic hexaoxazole exhibits significant cytotoxicity towards P388, RPMI 8402, and KB3-1 cell lines with IC₅₀ values of 45, 25, and 38 nM respectively.

The isolation and structure elucidation of telomestatin has sparked interest in the identification of compounds that can selectively stabilize G-quadruplex DNA.¹ These compounds have potential as a novel class of anticancer agents.² Recently the synthesis of the macrocyclic hexaoxazole HXDV was disclosed.³ HXDV is a 24-membered macrocycle that stabilizes G-quadruplexes presumably by π -stacking and hydrogen bonding interactions. HXDV is more readily synthesized than telomestatin. Despite the simpler structure, HXDV is extraordinarily selective at stabilizing G-quadruplex vs. duplex DNA, and is at least as potent as telomestatin as a cytotoxic agent.⁴ These results prompted us to investigate whether modifications could be made to the macrocyclic framework while retaining selectivity for G-quadruplex stabilization and antitumor activity. In this account a method is presented for the synthesis of macrocyclic hexaoxazoles *via* a ring closing metathesis reaction.

Previous studies in our laboratory confirmed that 24-membered macrocycles comprised of two teroxazole moieties joined by at least one valine residue displayed extraordinary levels of G-quadruplex stabilization with no detectable levels of duplex DNA stabilization.⁵ It was reasoned that two teroxazole units joined at their 2"- and 4-positions by a valine residue might be capable of undergoing a ring-closing reaction if the remaining 2"- and 4-positions were functionalized as allyl and vinyl respectively. Ring-closing metathesis has become a reliable method for the formation of a wide variety of cyclic compounds from five-membered rings to

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extremely large rings of 36–72 atoms.⁶ Certainly 24-membered rings have been synthesized previously by this method, although there is usually considerably more conformational flexibility in the acyclic precursors than in the present example.⁷ Recently, the application of the cross metathesis reaction to isolated vinyl oxazoles was reported.⁸ We report herein the first example of a ring-closing metathesis being applied to the synthesis of a macrocyclic hexaoxazole.

The synthesis of 2"-allylteroxazole-4-carboxylic acid is shown in Scheme 1. Teroxazole **1** is prepared in a similar manner as previously reported for **3**, but starting with oxazole-4-carboxylic acid.^{3,9} Deprotonation at the 2"-position was effected with LiHMDS although the resulting lithio derivative could not be directly alkylated with allyl bromide. Fortunately, the cuprate derived from 2"-lithio **1** did react with allyl bromide to give the 2"-allyl derivative in 37% yield. The *tert*-butyl group was removed to give **2** in 89% yield upon treatment with TFA at 0 °C.¹⁰

The preparation of 4-vinylteroxazole **5** (Scheme 2) was performed by routine transformation of the known **3**. Reduction of the ester with NaBH₄/LiCl followed by Swern oxidation afforded aldehyde **4**. A Wittig reaction of **4** with methylenetriphenylphosphorane followed by TFA catalyzed removal of the Boc group gave intermediate **5** in 54% yield for the two steps.¹⁰

2"-Allylteroxazole **2** and 4-vinylteroxazole **5** were coupled in 69% yield using EDC and HOBt in the absence of added base (Scheme 3). The allyl group was found to be extremely sensitive to isomerization to a 1-propenyl derivative in the presence of triethylamine or Hunig's base. Amide **6** was now set up to test the viability of the ring-closing metathesis reaction. The second generation Grubb's catalysts **7a** and the Hoveyda catalyst **7b** were evaluated for their ability to promote the macrocyclization (Figure 1).^{11,12} Both catalysts gave the desired macrocycle **8** although **7b** gave slightly higher yields (32% *vs.* 24% using **7a**). The reactions were remarkably clean with no evidence of any cross-metathesis by-products. The geometry of the newly-formed olefinic linkage is *E*- as determined from the vicinal coupling constant J = 15.6 Hz.

The unsaturated macrocycle **8** and HXDV were evaluated for their relative abilities to bind and thermally stabilize G-quadruplex and duplex DNA in the presence of K⁺ ions. We used salmon testes DNA (ST-DNA) as a representative model for duplex DNA. As a representative model for G-quadruplex DNA, we used the human telomeric DNA sequence $d[T_2G_3(T_2AG_3)_3A]$, which is known to form an intramolecular G-quadruplex in potassium solution. ¹³ Figure 2 shows the UV melting profiles (depicted in their first-derivative forms) of d $[T_2G_3(T_2AG_3)_3A]$ and ST-DNA in the absence and presence of the two compounds. Note that the presence of neither compound alters the thermal stability of ST-DNA, with any differences between the transition temperatures (T_{tran}) corresponding to the maxima of the first-derivative melting profiles being within the experimental uncertainty. This observation is indicative of an absence of duplex DNA binding on the part of both compounds. In the case of HXDV, we used isothermal titration calorimetry to verify the lack of duplex DNA binding implied by the UV melting studies.¹⁴

In striking contrast to their negligible impacts on ST-DNA thermal stability, both compounds increase the thermal stability of d[T₂G₃(T₂AG₃)₃A]. Thus, like HXDV, compound **8** binds to G-quadruplex DNA with a high degree of specificity. Significantly, the unsaturated macrocycle **8** thermally stabilizes d[T₂G₃(T₂AG₃)₃A] to a greater extent than HXDV ($\Delta T_{tran} = 24.5 \text{ °C}$ and 16.0 °C for compound **8** and HXDV, respectively). The greater ΔT_{tran} afforded by compound **8** relative to HXDV may reflect a correspondingly enhanced affinity for the host quadruplex, as the relative degree to which a ligand thermally stabilizes a nucleic acid target often correlates with its relative binding affinity.

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Evaluation of the cytotoxic activity of unsaturated macrocycle **8** was performed using an MTT assay (Table 1). Among the cell lines employed for this assay was murine leukemia P388, a human lymphoblastoma RPMI 8402, human nasopharyngeal carcinoma KB3-1, and KB3-1 cell lines that over-express for the efflux transporters MDR1 (KBV-1), and BCRP (KBH5.0). The data indicate that compound **8** displays potent cytotoxic activity against P388, RPMI 8402, and KB3-1 with IC₅₀ values of 45, 25, and 38 nM respectively. In each case the IC₅₀ values determined for compound **8** represents an order of magnitude improvement over those observed for the macrocyclic hexaoxazole HXDV. The data also suggest that both **8** and HXDV are substrates for MDR1, but not BCRP.

In summary, ring closing metathesis offers a rapid method for the synthesis of 24-membered macrocyclic hexaoxazoles from suitably substituted teroxazole precursors. The resulting unsaturated macrocycle **8** selectively stabilizes G-quadruplex DNA without any detectable stabilization of duplex DNA. The degree of G-quadruplex stabilization was greater than previously observed using HXDV. The replacement of one of the valine linkages of HXDV by a propenyl group led to an order of magnitude enhancement in cytotoxic potency when assayed against several tumor cell lines. These results when taken together suggest that the ring-closing metathesis approach has the potential for development of novel G-quadruplex stabilizers with enhanced anticancer activity.

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10. The structures of all compounds were determined using ¹H (200 MHz) and ¹³C-NMR (50 MHz) in CDCl₃ unless otherwise noted. Results are reported as ppm downfield from internal TMS: 1: white solid, mp 283 °C (dec.); ¹H NMR (CDCl₃) δ 8.44 (s, 2H), 8.23 (s, 1H), 8.07 (d, 1H, *J*=0.74), 1.61 (s, 9H); ¹³C NMR (CDCl₃) δ 159.1, 155.0, 151.1, 142.3, 138.8, 138.5, 135.0, 130.2, 128.8, 81.6, 27.3; IR (thin film, NaCl) 1721 cm⁻¹; HRMS (ESI) *m/z* calculated for C₁₄H₁₃N₃O₅Na (M+Na)

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326.0753; found 326.0750. 2: white solid, mp 283–285 °C (dec); IR (CHCl₃) 3135, 1686 cm⁻¹; ¹H NMR (CDCl₃+CD₃OD) δ 8.38 (s, 1H) 8.27 (s, 1H), 8.26 (s, 1H), 5.91 (m, 1H), 5.19 (m, 2H), 3.55 (dt, 2H, J = 1.4, 6.6 Hz); HRMS calcd for C₁₃H₉N₃O₅H (M+H): 288.0620; found: 288.0622. 4: white solid, mp 196–198 °C; IR (CHCl₃) 3373, 1690 cm⁻¹; ¹H NMR (CDCl₃) δ 10.0 (s, 1H), 8.38 (s, 1H) 8.33 (s, 1H), 8.32 (s, 1H), 5.29 (d, 1H, J = 9.2 Hz), 4.83 (dd, 1H, J = 6.2, 8.8 Hz), 2.23 (m, 1H), 1.42 (s, 9H), 0.94 (d, 3H, J = 4.0 Hz), 0.92 (d, 3H, J = 3.0 Hz); ¹³C NMR (CDCl₃) δ 183.1, 164.8, 155.5, 155.0, 154.4, 142.7, 140.7, 138.5, 138.4, 129.8, 128.7, 79.2, 53.4, 32.0, 27.4, 17.8, 17.1; HRMS calcd for C19H22N4O6Na (M+Na): 425.1437; found: 425.1432. 5: white solid, mp 125-127 °C; IR (CHCl₃) 3381 cm⁻¹; ¹H NMR (CDCl₃) δ 8.34 (s, 1H) 8.30 (s, 1H), 7.63 (s, 1H), 6.59 (dd, 1H, J = 11, 17.4 Hz), 6.06 (d, 1H, J = 17.4 Hz), 5.37 (d, 1H, J = 11 Hz), 3.92 (d, 1H, J = 6.6 Hz), 2.17 (m, 1H), 1.00 (d, 3H, J = 5.6 Hz), 0.96 (d, 3H, J = 7.0 Hz); ¹³C NMR (CDCl₃) δ 167.4, 155.2, 153.9, 139.6, 138.3, 137.4, 134.1, 130.7, 128.7, 123.8, 116.0, 55.0, 32.5, 18.1, 17.1; HRMS calcd for C₁₅H₁₆N₄O₃H (M+H): 301.1301; found: 301.1269. **6**: white solid, mp 235–237 °C (dec); IR (CHCl₃) 3363, 1653 cm-1; ¹H NMR (CDCl₃) δ 8.34 (s, 2H) 8.33 (s, 1H), 8.30 (s, 1H), 8.26 (s, 1H), 7.62 (s, 1H), 7.59 (d, 1H, J = 10.4 Hz), 6.57 (dd, 1H, J = 11, 17.4 Hz), 6.01 (m, 2H), 5.32 (m, 4H), 3.64 (d, 2H, *J* = 6.2 Hz), 2.41 (m, 1H), 1.07 (d, 3H, *J* = 7.0 Hz), 1.00 (d, 3H, *J* = 6.6 Hz); ¹³C NMR (CDCl₃) & 163.5, 163.3, 159.0, 155.6, 155.0, 153.9, 153.7, 140.7, 139.6, 138.6, 138.5, 138.2, 137.5, 135.9, 134.1, 130.7, 130.0, 129.5, 129.1, 128.8, 123.8, 118.3, 116.0, 51.6, 31.9, 31.6, 18.0, 17.5; HRMS calcd for C₂₈H₂₃N₇O₇H (M+H): 570.1737; found: 570.1732. 8: white solid, mp 195–197 ° C (dec); $[\alpha]_D = -53.8^{\circ}$ (c 0.433, CHCl₃); IR (CHCl₃) 3149, 1669, 1658 cm⁻¹; ¹H NMR (CD₂Cl₂, 400 MHz) δ 8.29 (d, 1H, J = 8.4 Hz), 8.258 (s, 1H) 8.254 (s, 1H), 8.23 (s, 1H), 8.20 (s, 1H), 8.18 (s, 1H), 7.60 (m, 1H), 7.56 (s, 1H), 6.49 (d, 1H, J = 15.6 Hz), 5.32 (m, 1H), 3.84 (ddd, 1H, J = 2.4, 2.8, 19.4 Hz), 3.73 (ddd, 1H, J = 8.4, 8.8, 20.2 Hz), 2.39 (m, 1H), 1.02 (d, 3H, J = 6.8 Hz), 0.94 (d, 3H, J = 6.8 Hz; ¹³C NMR (CDCl₃) δ 163.4, 163.1, 159.1, 155.7, 154.9, 153.9, 153.8, 140.1, 139.0, 137.6, 137.40, 137.38, 136.29, 136.27, 132.7, 130.8, 130.2, 129.0, 128.9, 127.7, 118.0, 52.2, 33.0, 28.8, 17.7, 17.1; HRMS calcd for C₂₆H₁₉N₇O₇H (M+H): 542.1424; found: 542.1429.

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Figure 2.

First derivatives of the UV melting profiles of $d[T_2G_3(T_2AG_3)_3A]$ (**A**) and ST-DNA (**B**) in the absence and presence of HXDV or compound **8**. Profiles were acquired on an AVIV model 14NT-UV-VIS spectrophotometer using quartz cuvettes with a 1 cm pathlength. The temperature was raised in 0.5 °C increments and the samples were allowed to equilibrate for 1.5 minutes at each temperature setting, whereupon absorbances were recorded over a period of 5 seconds and averaged. When present, ST-DNA was used at a base pair concentration of 15 µM, while $d[T_2G_3(T_2AG_3)_3A]$ was used at a strand concentration of 4 µM. Macrocyclic ligands were used at a concentration of 15 µM in the ST-DNA experiments and 20 µM in the $d[T_2G_3(T_2AG_3)_3A]$ experiments. The solution conditions were 10 mM EPPS (pH 7.5) and sufficient KCl to bring the total K⁺ concentration to 50 mM. In the ST-DNA experiments, the acquisition wavelength was 260 nm, while being 295 nm in the $d[T_2G_3(T_2AG_3)_3A]$



Scheme 1.

(a) LiHMDS, THF, -42 °C; CuI, allyl bromide, -42 °C to rt, 37%; (b) TFA, anisole/CH₂Cl₂ (1:5), 0 °C, 89%.

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(a) NaBH₄, LiCl, THF/MeOH (1:1) 73 %; (b) oxalyl chloride, DMSO, CH₂Cl₂, Et₃N, -78 °C, 88%; (c) Ph₃PCH₃Br, n-BuLi, THF, 0 °C to rt, 74%; (d) TFA, CH₂Cl₂, 0 °C, 73%.



Scheme 3.

(a) EDC, HOBt, DMF, 0 °C to rt, 69%; (b) **7b**, CH_2Cl_2 , Δ , 96 h, 32%. Structure of HXDV shown for comparison.

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KBH5.0	0.043 0.5
KBV-1	3.3 8.0
KB3-1	0.038 0.3
RPMI 8402	0.025 0.41
P388	0.045 0.5
Compound	8 HXDV

^aIC50 values are reported in \square M concentrations and represent the average values from replicate assays. The exposure time of the cells to the test compounds was 4 days.