

# G-Quartet Formation from an N<sup>2</sup>-Modified Guanosine Derivative

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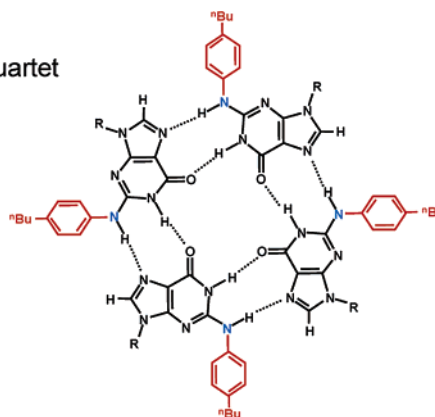
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## ABSTRACT

### An N<sup>2</sup>-modified G-quartet



We report G-quartet formation from an N<sup>2</sup>-modified lipophilic guanosine nucleoside, N<sup>2</sup>-(4-*n*-butylphenyl)-2',3',5'-*O*-triacetylguanosine. We show that, in the presence of either K<sup>+</sup> or Na<sup>+</sup>, this guanosine derivative self-assembles into a D<sub>4</sub>-symmetric octamer consisting of two stacking all-*syn* G-quartets in a tail-to-tail (or head-to-head) fashion and a central ion.

G-Quartet formation is the fundamental driving force for a large number of guanosine derivatives including G-rich DNA and RNA oligomers to form highly ordered structures.<sup>1–5</sup> The G-quartet motif has not only been found in important regions in human genome such as telomeres,<sup>6,7</sup> but also shown potential usefulness in nanotechnology.<sup>8,9</sup> However, only a few cases are known to date where base-modified guanosine nucleosides are involved in G-quartet formation,

and the modifications occur exclusively at the C<sub>8</sub> position of the guanine.<sup>10–13</sup> There are also two studies reporting improved G-quadruplex stability from N<sup>2</sup>-modified DNA oligomers.<sup>14,15</sup> Here we report the first example of G-quartet formation from an N<sup>2</sup>-modified lipophilic guanosine nucleoside: N<sup>2</sup>-(4-*n*-butylphenyl)-2',3',5'-*O*-triacetylguanosine (G1); see Figure 1.<sup>16</sup>

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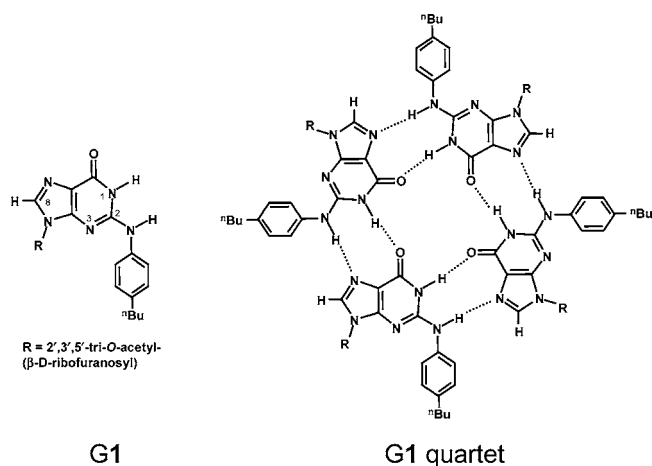
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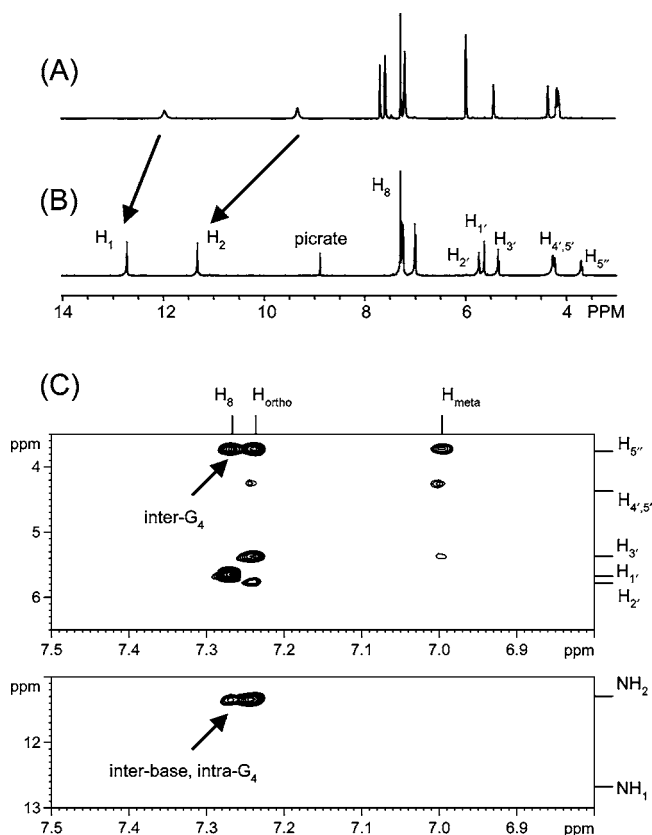
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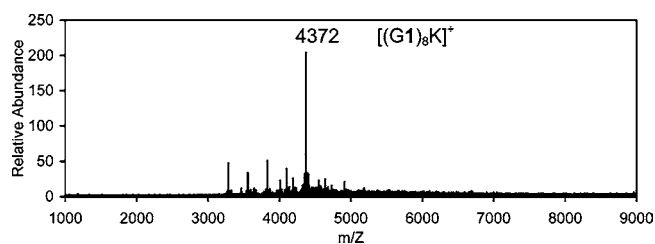


**Figure 1.** Molecular structure of an  $N^2$ -modified guanosine and the G-quartet formation.

Figure 2 shows the  $^1\text{H}$  NMR spectra of **G1** in both monomeric and aggregate forms in  $\text{CDCl}_3$ .<sup>17</sup> The imino



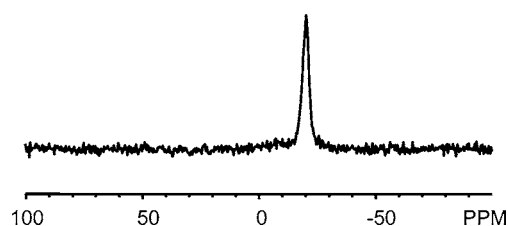
**Figure 2.** Portions of 1D  $^1\text{H}$  NMR spectra for (A) **G1** monomer and (B) **G1** aggregate (in the presence of  $\text{K}^+$  picrate) in  $\text{CDCl}_3$  at 298 K. (C) A region of the 2D NOESY spectrum of **G1** aggregate at 298 K. A mixing time of 300 ms was used. A total of 64 transients were collected for each of the 256  $t_1$  increments with a recycle delay of 2 s. All  $^1\text{H}$  NMR spectra were obtained on a Bruker Avance-600 spectrometer ( $B_0 = 14.1$  T).



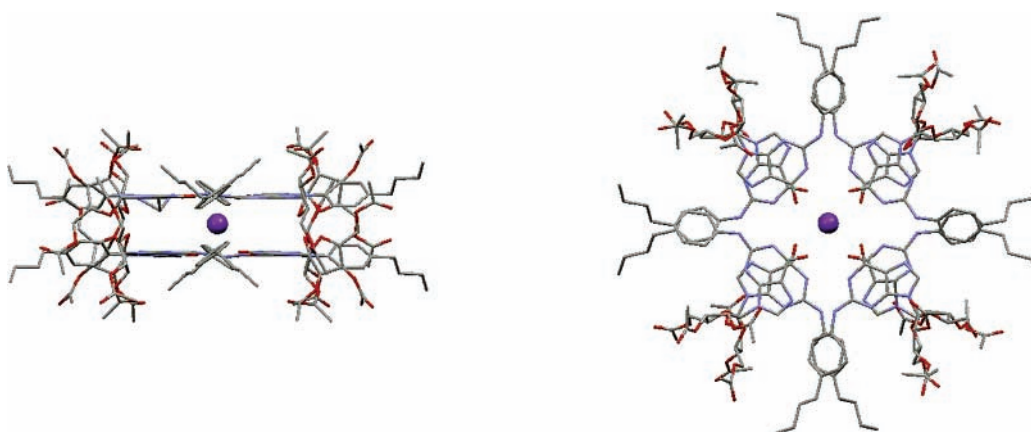
**Figure 3.** Positive-ion electrospray mass spectrum of **G1** aggregate prepared by liquid–liquid extraction of  $\text{K}^+$  picrate into chloroform containing **G1**. The theoretical molecular weight for  $[(\text{G1})_8\text{K}]^+$  is 4364.5.

( $N^1\text{H}_1$ ) and amino ( $N^2\text{H}_2$ ) signals for **G1** monomer (Figure 2A) are shifted considerably toward the high-frequency direction when interacting with  $\text{K}^+$  picrate, characteristic of G-quartet formation (Figure 2B).<sup>5</sup> As seen in Figure 2C, an inter-base NOE cross-peak is observed between  $\text{H}_8$  and  $\text{NH}_2$  signals, which is also a spectral signature for G-quartet formation.<sup>5</sup> In addition, strong cross-peaks are observed between  $\text{H}_8$  and  $\text{H}_{1'}$  and between phenyl and ribose protons. These are indicative of a *syn* conformation with respect to the  $\text{N}_9\text{--C}_{1'}$  glycosidic bond. The crystal structure of **G1** monomer (crystallized from  $\text{CH}_3\text{CN}$  in the absence of alkali metal ions) shows that **G1** adopts a *syn* conformation ( $\chi_{\text{CN}} = 75.7^\circ$ ).<sup>18</sup>

Because only one set of  $^1\text{H}$  NMR signals were observed for the **G1** aggregate, all **G1** molecules must adopt a *syn* conformation. The  $^1\text{H}$  NMR signal integrations yield a **G1**/picrate ratio of 8:1, suggesting the formation of a **G1** octamer. To verify the size of the **G1** aggregate, we used a pulse-field-gradient diffusion NMR technique.<sup>19</sup> The translational diffusion coefficients determined for the **G1** monomer and aggregate in  $\text{CDCl}_3$  at 298.2 K are  $6.06 \pm 0.06 \times 10^{-10}$  and  $3.09 \pm 0.06 \times 10^{-10} \text{ m}^2 \text{ s}^{-1}$ , respectively. This observed  $D_{\text{8mer}}/D_{\text{monomer}}$  ratio of 0.51 is consistent with the formation of a **G1** octamer.<sup>20,21</sup> As shown in Figure 3, the positive-ion electrospray mass spectrum of the **G1** aggregate confirms that the octamer formation is predominant. Within the octamer, the two G-quartets are likely twisted by 30–45°



**Figure 4.**  $^{23}\text{Na}$  magic-angle-spinning NMR spectrum of **G1** prepared in the presence of  $\text{NaClO}_4$ . Solid-state  $^{23}\text{Na}$  NMR spectra were obtained on a Bruker Avance-500 spectrometer using a 4-mm double-resonance MAS probe. Experimental parameters: 9 kHz spinning, 7458 transients, 2 s recycle delay, and 70 kHz proton decoupling.



**Figure 5.** Molecular model (side and top views) for a  $D_4$ -symmetric G1 octamer,  $[(G1)_8K^+]$ , where the two G-quartets are oriented in a tail-to-tail fashion. Hydrogen atoms are omitted for clarity.

with respect to each other, as supported by the fact that an inter-quartet NOE cross-peak is observed between  $H_8$  and  $H_5''$  signals.

To probe the alkali metal ion binding site, we obtained a solid-state  $^{23}Na$  NMR spectrum for the G1 aggregate prepared in the presence of  $Na^+$  ions. As shown in Figure 4, a relatively narrow signal centered at  $-19$  ppm was observed in the magic-angle spinning  $^{23}Na$  NMR spectrum, suggesting that the  $Na^+$  ion is sandwiched between two G-quartets, similar to the situations found in other G-quartet structures including DNA oligomers.<sup>22–24</sup>

Based on the experimental data, we can conclude that, in the presence of  $K^+$  or  $Na^+$ , G1 self-associates into two

stacking all-*syn* G-quartets in a tail-to-tail (or head-to-head) arrangement with a central metal ion forming a  $D_4$ -symmetric G1 octamer, as illustrated in Figure 5. This represents another example for this very rare type of (iso)guanine octamer.<sup>21,25</sup> The molecular model shown in Figure 5 also suggests a possible  $\pi$ - $\pi$  stacking between the phenyl rings from the two different G-quartets. The observed  $^1H$  chemical shift changes for the ring protons between monomeric and aggregate forms ( $\Delta\delta = 0.34$  and  $0.18$  ppm for the *ortho*

(16) G1 was synthesized by following the literature procedure: Wright, G. E.; Dudycz, Z. W. *J. Med. Chem.* **1984**, *27*, 175–181.

(17) Preparation of G1 aggregate was achieved by either liquid–liquid extraction or solid–liquid extraction. See: Forman, S. L.; Fettingner, J. C.; Pieraccini, S.; Gottarelli, G.; Davis, J. T. *J. Am. Chem. Soc.* **2000**, *122*, 4060–4067.

(18) Single crystals of G1 monomer were obtained from slow evaporation of a  $CH_3CN$  solution in the absence of alkali salts. The crystal data were collected on a Bruker P4 single-crystal X-ray diffractometer with a Smart CCD-1000 detector, Mo  $K\alpha$  radiation, (50 kV and 30 mA) at 180 K. The structure was solved by using the direct method. Crystal data: formula  $C_{26}H_{30}N_5O_8$ ,  $a = 9.019(2)$ ,  $b = 10.726(3)$ ,  $c = 14.449(4)$  Å,  $\beta = 105.111(4)^\circ$ ,  $V = 1349.5(6)$  Å<sup>3</sup>,  $Z = 2$ , monoclinic space group  $P2_1$ . Convergence to  $R_1 = 0.055$ ,  $wR_2 = 0.126$  ( $I > 2\sigma(I)$ ), GOF = 0.942 was achieved by using 4735 reflections and 353 parameters. All non-hydrogen atoms were refined anisotropically. Details of the crystal structure can be found in the supporting materials.

(19) All  $^1H$  diffusion NMR spectra were recorded on a Bruker Avance-600 spectrometer using a pulse sequence of longitudinal-eddy-current delay (LED) with bipolar-gradient pulses. The pulse field gradient duration was varied from 4 to 15 ms, and the variable gradient strength ( $G$ ) was changed from 6 to 350 mT/m. The diffusion period was varied from 50 to 90 ms. A total of 16 transients were collected for each of the 16 or 32 increment steps with a recycle delay of 20 s. The eddy-current delay was set to 5  $\mu$ s.

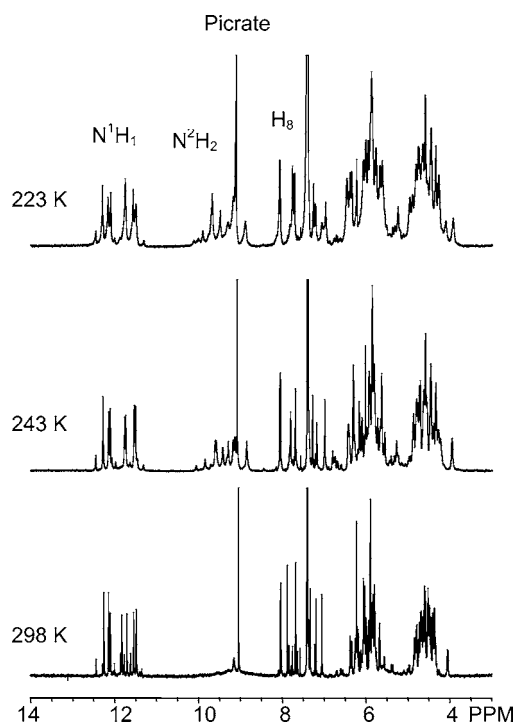
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**Figure 6.** Portions of 1D  $^1H$  NMR spectra for G2 aggregate prepared by liquid–liquid extraction of  $K^+$  picrate into chloroform solution of G2.

and *meta*  $^1\text{H}$  resonances, respectively) are consistent with the formation of  $\pi$ - $\pi$  stacking.

To further examine the consequence of  $\text{N}^2$ -alkylation, we compared the G-quartet formation from unmodified 2',3',5'-*O*-triacetylguanosine (**G2**). As seen from Figure 6, the  $^1\text{H}$  NMR spectra for **G2** aggregates in the presence of  $\text{K}^+$  and  $\text{Na}^+$  ions exhibit very complex features, indicating the formation of polymeric columnar aggregates. The 2D  $^1\text{H}$  NOESY spectra confirmed the formation of G-quartet in these systems. The ESI-MS spectra for **G2**/ $\text{K}^+$  aggregate show clearly the presence of octamer, dodecamer, and hexadecamer; see Supporting Information. Thus, the  $\text{N}^2$ -alkylation in **G1** changes the G-quadruplex structure from an extended polymer to a discrete octamer. This increased stability of the **G1** octamer formation is presumably due to the fact that the  $\text{N}^2$ -alkylation stabilizes the *syn* conformation. It is also possible that the  $\pi$ - $\pi$  stacking between the phenyl rings further favors the formation of a tail-to-tail (or head-to-head) stacking, resulting in a discrete  $D_4$ -symmetric octamer.

Although the present study is concerned with G-quartet formation from a lipophilic guanosine nucleoside in organic solvent, our finding that  $\text{N}^2$ -substitution does not hinder G-quartet formation may have implications in the design of new G-quadruplex structures formed by DNA and RNA. In this regard, there are several known cases where the exocyclic amino hydrogen atom that is not involved in the G-quartet formation participates in additional intermolecular hydrogen bonding in DNA and RNA G-quadruplexes.<sup>26–30</sup> These studies suggest that recognition of the  $\text{N}^3$ - $\text{N}^2$  edge

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of the G-quartet, which is located at the groove region of a G-quadruplex, may be important for DNA and RNA folding. Consequently, any modification at the  $\text{N}^2$  site of the guanine base may interfere with nucleic acid folding and recognition. We anticipate that residue-specific  $\text{N}^2$  modification can be exploited as a new handle for fine-tuning G-quadruplex structure and function.

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**Note added in proof.** After submission of this work, we have learned that Kaucher and Davis also observed G-quartet formation from an  $\text{N}^2$ ,  $\text{C}_8$ -disubstituted guanosine derivative.<sup>31</sup>

**Supporting Information Available:** COSY and NOESY spectra for **G1** aggregates and a table containing  $^1\text{H}$  chemical shifts; complete crystal data (in CIF format); diagrams showing the structure of **G1** and NOESY and ESI-MS spectra for **G2**/ $\text{K}^+$  aggregates. This material is available free of charge via the Internet at <http://pubs.acs.org>.

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