

# Direct $^{23}\text{Na}$ NMR observation of mixed cations residing inside a G-quadruplex channel†

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We report direct  $^{23}\text{Na}$  NMR observation for the presence of mixed cations ( $\text{Na}^+/\text{K}^+$ ,  $\text{Na}^+/\text{Rb}^+$ ,  $\text{Na}^+/\text{Sr}^{2+}$ ) inside the G-quadruplex channel formed by the self-association of guanosine 5'-monophosphate at pH 8.

Alkali metal cations such as  $\text{Na}^+$  and  $\text{K}^+$  are known to play an important role in stabilizing G-quadruplex structures.<sup>1</sup> Although solid-state techniques such as X-ray crystallography and solid-state NMR are quite useful for localizing alkali metal cations in G-quadruplex DNA,<sup>2,3</sup> it is highly desirable to have biophysical techniques that can detect these cations in solution. To this end, NMR methodologies based on spin-1/2 probes such as  $^{15}\text{NH}_4^+$  and  $^{205}\text{Tl}^+$  have been developed and successfully applied to G-quadruplex DNA.<sup>4,5</sup> Recently, we demonstrated that solution metal ( $^{23}\text{Na}$ ,  $^{39}\text{K}$ ,  $^{87}\text{Rb}$ ) NMR can be used for direct detection of alkali metal cations in G-quadruplex DNA.<sup>6</sup> Here we report that high-resolution  $^{23}\text{Na}$  NMR spectra allow direct detection of mixed cations residing inside a G-quadruplex channel.

Fig. 1 shows a  $^{23}\text{Na}$  NMR spectrum for  $\text{Na}_2(5'\text{-GMP})$  at pH 8.† This spectrum exhibits two peaks, one at  $\delta(^{23}\text{Na})$  0 ppm having

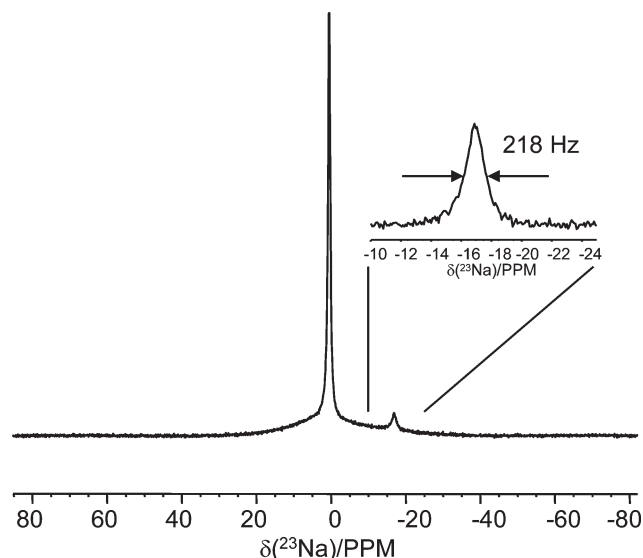


Fig. 1  $^{23}\text{Na}$  NMR spectrum for 1.0 M  $\text{Na}_2(5'\text{-GMP})$  at 5 °C.

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a bi-Lorentzian line shape and the other at  $\delta(^{23}\text{Na})$  -17 ppm. As we have shown recently,<sup>6</sup> the former signal is due to free  $\text{Na}^+$  ions and the latter is assigned to  $\text{Na}^+$  ions residing inside the G-quadruplex channel. At pH 8,  $\text{Na}_2(5'\text{-GMP})$  is known to self-associate into molecular cylinders of 10–30 nm in length, depending on the actual 5'-GMP concentration.<sup>7</sup> Under such a circumstance, the  $\text{Na}^+$  ions residing inside the G-quadruplex channel can be seen as tightly bound to a macromolecule with an effective molecular weight of 30–100 kDa. An additional benefit of  $\text{Na}^+$  binding to this large molecular species is that the line width for the slow decaying component becomes narrow again when the spin-3/2 system is far from the so-called extreme narrowing condition.<sup>8</sup> In the present case, the full width at the half height (FWHM) of the channel  $^{23}\text{Na}$  NMR signal is only 218 Hz (<1.4 ppm at 14.1 T) at 5 °C. With such a high spectral resolution, it may be possible that the  $^{23}\text{Na}$  chemical shift can be used as a sensitive reporter for the nature of cations occupying the neighboring site inside a G-quadruplex channel. Indeed, the existence of G-quadruplexes containing mixed cations has been observed recently in  $^1\text{H}/^{15}\text{N}$  NMR and X-ray crystallographic studies.<sup>9</sup>

Fig. 2 shows portions of the  $^{23}\text{Na}$  NMR spectra for 5'-GMP containing mixed cations  $\text{Na}^+/\text{M}^{n+}$  ( $\text{M}^{n+} = \text{K}^+, \text{Rb}^+, \text{Sr}^{2+}$ ). It is

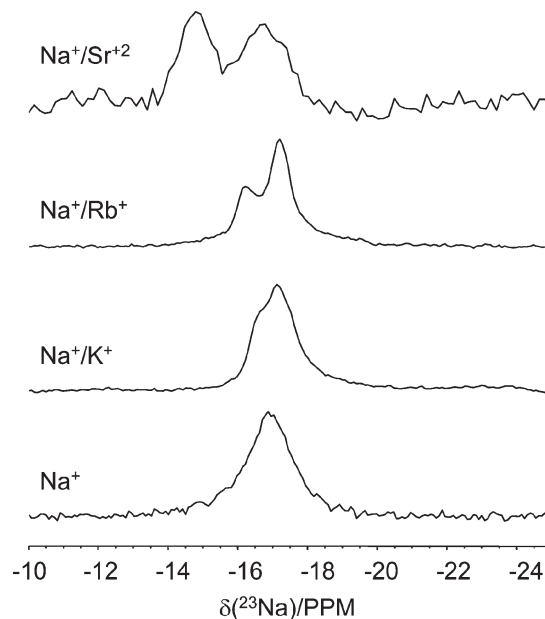
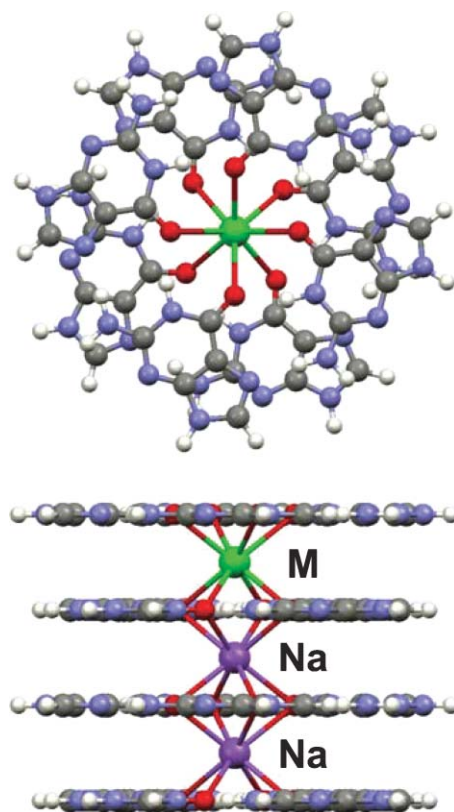


Fig. 2 Portions of the  $^{23}\text{Na}$  NMR spectra of 1.0 M  $\text{Na}_2(5'\text{-GMP})$  containing mixed cations at 5 °C. The concentrations of the added cations are:  $\text{K}^+$ , 100 mM;  $\text{Rb}^+$ , 100 mM;  $\text{Sr}^{2+}$ , 20 mM.

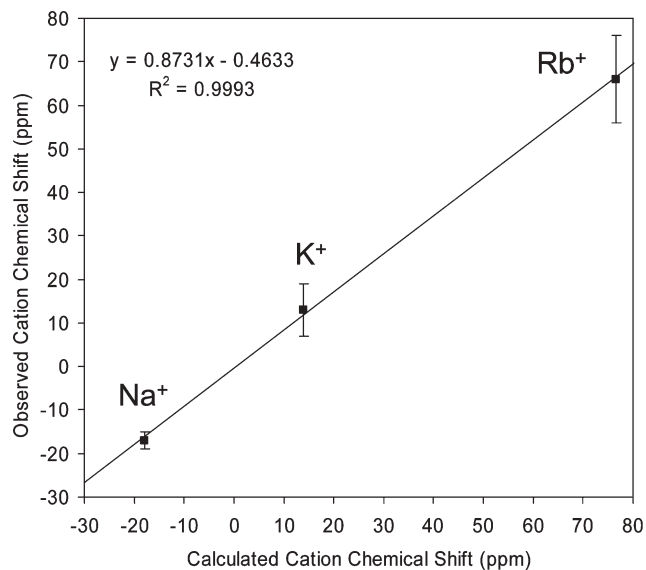
well known that  $K^+$ ,  $Rb^+$  and  $Sr^{2+}$  are all capable of entering a G-quadruplex channel.<sup>1</sup> As seen in Fig. 2, when  $M^{n+}$  is added to  $Na_2(5'-GMP)$  solution, a new signal appears in the  $^{23}Na$  NMR spectra. In the cases studied here, each of the new signals exhibits a less negative  $^{23}Na$  chemical shift (corresponding to a less shielded environment) than that of the original  $^{23}Na$  signal. We assign these new signals to the  $Na^+$  ions with one of the two neighboring sites being occupied by  $M^{n+}$ , *i.e.*  $G_4-M-G_4-Na-G_4-Na-G_4$ . Because the concentration of the added cations employed in our study is always much smaller than that of  $Na^+$ , we can safely neglect the population of the  $Na^+$  ions with both neighboring sites being occupied by  $M^{n+}$  ions, *i.e.*  $G_4-M-G_4-Na-G_4-M-G_4$ . We also performed a Na/Rb titration experiment in which various amounts of  $Rb^+$  ions were added to the  $Na_2(5'-GMP)$  solution. As expected, as the concentration of  $Rb^+$  ions increases, the relative intensity of the  $^{23}Na$  signal at  $\delta(^{23}Na) - 16.2$  ppm also increases, indicating that more  $Rb^+$  ions have entered the channel (see ESI†).

Another important point worth emphasizing is that the observed  $^{23}Na$  NMR spectra will provide clues to the rate of cation movement through the channel. The fact that a separate  $^{23}Na$  NMR signal is actually observed for the channel  $Na^+$  ions with  $M^{n+}$  as an immediate neighbor immediately suggests that the movement of the added  $M^{n+}$  ions ( $K^+$ ,  $Rb^+$  and  $Sr^{2+}$ ) in and out of the G-quadruplex channel must be slow on the  $^{23}Na$  NMR time scale employed in this study. In particular, the smallest signal separation observed for the different channel  $Na^+$  ions is 0.6 ppm (corresponding to 95 Hz at 14.1 T) as observed in the Na/K case. This indicates that the averaged residence time of  $K^+$  ions inside the channel (not necessarily the residence time at a particular site) must be much longer than  $(2\pi \times 95 \text{ Hz})^{-1} \approx 2$  ms. Otherwise, the two  $^{23}Na$  NMR signals would be averaged into one. This is consistent with recent findings that the residence time of  $NH_4^+$  ions in a G-quadruplex channel containing mixed  $Na^+/NH_4^+$  ions is 36 ms<sup>8</sup> and that the residence time of  $Tl^+$  ions in  $d(G_4T_4G_4)$  is approximately 100 ms.<sup>5</sup>

To further verify our  $^{23}Na$  NMR spectral assignment, we performed *ab initio*  $^{23}Na$  chemical shielding calculations for  $Na^+$  ions inside a G-quadruplex channel. Our model shown in Fig. 3 consists of four stacking G-quartets and three channel cations (a total of 259 atoms). Each cation is sandwiched between two adjacent G-quartets that are separated by 3.4 Å and twisted by 45°. To model the mixed cation cases, the top  $Na^+$  ion is replaced by  $K^+$ ,  $Rb^+$  and  $Sr^{2+}$ , respectively, and the magnetic shielding at the central  $Na^+$  ion is calculated. The computational results shown in Table 1 confirm that, when a  $K^+$  ion (or  $Rb^+$  and  $Sr^{2+}$ ) occupies the neighboring channel site, the central  $Na^+$  ion experiences a slightly less shielding environment. Furthermore, the observed trend on going from Na, K, Rb to Sr is well reproduced by quantum chemical calculations. Considering the approximations used in the model, the agreement between the experimental and calculated  $^{23}Na$  chemical shifts is remarkable. To rule out the possibility of a  $Na^+$  binding within the G-quartet plane, we performed further shielding calculations for a  $G_4-Na^+$  model where the  $Na^+$  ion is located at the center (in-plane) of the G-quartet. The calculations with different basis sets consistently predict that the  $^{23}Na$  chemical shift for such a  $Na^+$  ion should be approximately +6 ppm. We have never observed any  $^{23}Na$  signal at this chemical shift for our 5'-GMP samples. Another way to check the validity of our model is to calculate the chemical



**Fig. 3** Top (upper) and side (lower) views of the G-quadruplex model used for chemical shielding calculations where  $M = Na, K, Rb$  and  $Sr$ .



**Fig. 4** Comparison between calculated and observed chemical shifts for  $Na^+$ ,  $K^+$ , and  $Rb^+$  ions residing inside the G-quadruplex channel.

**Table 1** Observed and calculated  $^{23}Na$  chemical shifts (in ppm) for  $Na^+$  ions inside a G-quadruplex containing mixed cations<sup>a</sup>

|            | $Na^+$ | $K^+$ | $Rb^+$ | $Sr^{2+}$ |
|------------|--------|-------|--------|-----------|
| Observed   | -17.0  | -16.6 | -16.2  | -14.8     |
| Calculated | -18.0  | -17.5 | -17.6  | -14.8     |

<sup>a</sup> See footnote for computational details.

shielding values at the  $K^+$  and  $Rb^+$  sites using the same model, because  $^{39}K$  and  $^{87}Rb$  NMR signatures for these cations residing between two G-quartets have also been established.<sup>6</sup> As seen in Fig. 4, the calculated chemical shifts are in good agreement with the experimental values. This provides strong evidence that the G-quadruplex model used in our calculations is reasonable.

In summary, our  $^{23}Na$  NMR results illustrate the remarkable resolution achievable in solution-state  $^{23}Na$  NMR spectra for  $Na^+$  ions tightly bound to a large molecular self-assembly. It is anticipated that similarly high resolution should be observed for  $Na^+$  ions bound to other biological macromolecules. Possible extensions of this type of experiment to  $^{39}K$  and  $^{87}Rb$  NMR are under way in our laboratory.

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## Notes and references

‡ The hydrated disodium salt of 5'-GMP (>99% purity) was purchased from Sigma-Aldrich. All  $^{23}Na$  NMR spectra were obtained on a Bruker Avance-600 NMR spectrometer operating at 600.13 and 158.76 MHz for  $^1H$  and  $^{23}Na$  nuclei, respectively. All  $^{23}Na$  chemical shifts are referenced to  $Na^+(aq.)$  at  $\delta = 0$  ppm.

§ Quantum chemical calculations were performed using the Gaussian 03 suite of programs<sup>10</sup> on a SunFire 6800 symmetric multiprocessor system. Each of the four nodes is equipped with  $24 \times 1.05$  GHz (8 MB E-Cache) UltraSPARC-III processor and 96 GB of RAM. For the central Na, a high-level correlation consistent basis set, cc-pV5Z, was used. For the two outer metal cations (Na, K, Rb and Sr), the all-electron pVTZ basis sets of Sadlej<sup>11</sup> were used. A 3-21G(d) basis set was used for all other non-metal atoms. Shielding calculations were performed at the Hartree-Fock (HF) level using the GIAO method as implemented in Gaussian 03. The computed absolute shielding ( $\sigma$ ) was converted to the chemical shift ( $\delta$ ) scale using  $\delta = \sigma_{ref} - \sigma$ , where  $\sigma_{ref}$  is the absolute shielding constant for the reference sample,  $Na^+(aq.)$ . We used  $\sigma_{ref} = 587.6$  ppm, which is the value calculated for  $[Na(H_2O)_6]^+$  (Na-O<sub>w</sub>: 2.433 Å) at the HF/6-31G(d)/cc-pVQZ level.<sup>12</sup> For calculations of  $^{39}K$  and  $^{87}Rb$  chemical shifts, we used  $\sigma_{ref} = 1246$  and 3218 ppm, respectively. These values were calculated at the HF/3-21G(d)/pVTZ levels for fully hydrated clusters,  $[K(H_2O)_8]^+$  (K-O<sub>w</sub>: 2.712 Å) and  $[Rb(H_2O)_8]^+$  (Rb-O<sub>w</sub>: 3.000 Å). All computations were performed at the High Performance Computing Virtual Laboratory (HPCVL) at Queen's University. Each calculation takes about 3–4 days of CPU time.

1 See reviews: W. Guschlbauer, J.-F. Chantot and D. Thiele, *J. Biomol. Struct. Dyn.*, 1990, **8**, 491–511; D. Sen and W. Gilbert, *Methods Enzymol.*, 1992, **211**, 191–199; J. R. Williamson, *Annu. Rev. Biophys. Biomol. Struct.*, 1994, **23**, 703–730; D. E. Gilbert and J. Feigon, *Curr. Opin. Struct. Biol.*, 1999, **9**, 305–314; M. A. Keniry, *Biopolymers*, 2001, **56**, 123–146; S. Neidle and G. N. Parkinson, *Curr. Opin. Struct. Biol.*, 2003, **13**, 275–283; J. T. Davis, *Angew. Chem., Int. Ed.*, 2004, **43**, 668–698.

- 2 G. Laughlan, A. I. H. Murchie, D. G. Norman, M. H. Moore, P. C. E. Moody, D. M. J. Lilley and B. Luisi, *Science*, 1994, **265**, 520–524; K. Phillips, Z. Dauter, A. I. H. Murchie, D. M. J. Lilley and B. Luisi, *J. Mol. Biol.*, 1997, **273**, 171–182; M. P. Horvath and S. C. Schultz, *J. Mol. Biol.*, 2001, **310**, 367–377; S. Haider, G. N. Parkinson and S. Neidle, *J. Mol. Biol.*, 2002, **320**, 189–200; G. N. Parkinson, M. P. H. Lee and S. Neidle, *Nature*, 2002, **417**, 876–880; P. Hazel, G. N. Parkinson and S. Neidle, *J. Am. Chem. Soc.*, 2006, **128**, 5480–5487.
- 3 D. Rovnyak, M. Baldus, G. Wu, N. V. Hud, J. Feigon and R. G. Griffin, *J. Am. Chem. Soc.*, 2000, **122**, 11423–11429; G. Wu and A. Wong, *Chem. Commun.*, 2001, 2658–2659; A. Wong, J. C. Fettinger, S. L. Forman, J. T. Davis and G. Wu, *J. Am. Chem. Soc.*, 2002, **124**, 742–743; A. Wong and G. Wu, *J. Am. Chem. Soc.*, 2003, **125**, 13895–13905; G. Wu, A. Wong, Z. Gan and J. T. Davis, *J. Am. Chem. Soc.*, 2003, **125**, 7182–7183; G. Wu and A. Wong, *Biochem. Biophys. Res. Commun.*, 2004, **323**, 1139–1144; R. Ida and G. Wu, *Chem. Commun.*, 2005, 4294–4296.
- 4 N. V. Hud, P. Schultze and J. Feigon, *J. Am. Chem. Soc.*, 1998, **120**, 6403–6404; N. V. Hud, P. Schultze, V. Sklenář and J. Feigon, *J. Mol. Biol.*, 1999, **285**, 233–243.
- 5 S. Bazu, A. A. Szewczak, M. Cocco and S. A. Strobel, *J. Am. Chem. Soc.*, 2000, **122**, 3240–3241; M. L. Gill, S. A. Strobel and J. P. Loria, *J. Am. Chem. Soc.*, 2005, **127**, 16723–16732.
- 6 A. Wong, R. Ida and G. Wu, *Biochem. Biophys. Res. Commun.*, 2005, **337**, 363–366.
- 7 A. Wong, R. Ida, L. Spindler and G. Wu, *J. Am. Chem. Soc.*, 2005, **127**, 6990–6998.
- 8 P. S. Hubbard, *J. Chem. Phys.*, 1970, **53**, 985–987; T. E. Bull, *J. Magn. Reson.*, 1972, **8**, 344–353; A. Delville, C. Detellier and P. Laszlo, *J. Magn. Reson.*, 1979, **34**, 301–315.
- 9 N. V. Hud, F. W. Smith, F. A. L. Anet and J. Feigon, *Biochemistry*, 1996, **35**, 15383–15390; C. Caceres, G. Wright, C. Gouyette, G. Parkinson and J. A. Subirana, *Nucleic Acids Res.*, 2004, **32**, 1097–1102; B. Pan, Y. Xiong, K. Shi, J. Deng and M. Sundaralingam, *Structure*, 2003, **11**, 815–823; P. Šket, M. Črnugelj, W. Koźmiński and J. Plavec, *Org. Biomol. Chem.*, 2004, **2**, 1970–1973; P. Šket, M. Črnugelj and J. Plavec, *Nucleic Acids Res.*, 2005, **33**, 3691–3697.
- 10 Gaussian 03, Revision C.02, M. J. Frisch, G. W. Trucks, H. B. Schlegel, G. E. Scuseria, M. A. Robb, J. R. Cheeseman, J. A. Montgomery, Jr., T. Vreven, K. N. Kudin, J. C. Burant, J. M. Millam, S. S. Iyengar, J. Tomasi, V. Barone, B. Mennucci, M. Cossi, G. Scalmani, N. Rega, G. A. Petersson, H. Nakatsuji, M. Hada, M. Ehara, K. Toyota, R. Fukuda, J. Hasegawa, M. Ishida, T. Nakajima, Y. Honda, O. Kitao, H. Nakai, M. Klene, X. Li, J. E. Knox, H. P. Hratchian, J. B. Cross, V. Bakken, C. Adamo, J. Jaramillo, R. Gomperts, R. E. Stratmann, O. Yazyev, A. J. Austin, R. Cammi, C. Pomelli, J. W. Ochterski, P. Y. Ayala, K. Morokuma, G. A. Voth, P. Salvador, J. J. Dannenberg, V. G. Zakrzewski, S. Dapprich, A. D. Daniels, M. C. Strain, O. Farkas, D. K. Malick, A. D. Rabuck, K. Raghavachari, J. B. Foresman, J. V. Ortiz, Q. Cui, A. G. Baboul, S. Clifford, J. Cioslowski, B. B. Stefanov, G. Liu, A. Liashenko, P. Piskorz, I. Komaromi, R. L. Martin, D. J. Fox, T. Keith, M. A. Al-Laham, C. Y. Peng, A. Nanayakkara, M. Challacombe, P. M. W. Gill, B. Johnson, W. Chen, M. W. Wong, C. Gonzalez and J. A. Pople, Gaussian, Inc., Wallingford, CT, 2004.
- 11 A. J. Sadlej, *Theor. Chim. Acta*, 1992, **81**, 45–63.
- 12 A. Wong, R. D. Whitehead, Z. Gan and G. Wu, *J. Phys. Chem. A*, 2004, **108**, 10551–10559.