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Direct Detection of Potassium Cations Bound to G-Quadruplex Structures by Solid-State 39K NMR at 19.6 T

Gang Wu,*,† Alan Wong,† Zhehong Gan,‡ and Jeffery T. Davis§

Department of Chemistry, Oueen's University, Kingston, Ontario, Canada K7L 3N6. Center of Interdisciplinary Magnetic Resonance, National High Magnetic Field Laboratory, 1800 East Paul Dirac Drive, Tallahassee, Florida 32310, and Department of Chemistry and Biochemistry, University of Maryland, College Park, Maryland 20742

Received January 7, 2003; E-mail: gangwu@chem.queensu.ca

The presence of K⁺ ions in living cells is believed to be crucial for the stability of G-quadruplex structures found in telomeric DNA and other G-rich sequences. 1-5 Recent crystallographic studies have yielded reliable information about the K⁺ ion coordination geometry in G-quadruplexes,6-9 confirming an earlier proposal that the K+ ion is sandwiched between two G-quartets. 10 Because of the critical role that monovalent cations play in G-quadruplex formation, considerable efforts have also been devoted to the development of other spectroscopic techniques for detecting cation binding in G-quadruplexes. Successful NMR applications have been demonstrated to use 23 Na (spin $^{3}/_{2}$), 15 N (spin $^{1}/_{2}$), and 205 Tl (spin $^{1}/_{2}$) as NMR probes to directly study Na⁺, NH₄⁺, and Tl⁺ ions in G-quadruplexes.^{11–16} Smirnov et al. recently showed that extended X-ray absorption fine structure (EXAFS) can be used to characterize the Pb²⁺ binding site in G-quadruplexes.¹⁷ In general, the rather weak association between K+ ions and biological structures renders solution ³⁹K (spin ³/₂) NMR spectroscopy to be of limited utility. ^{10,18} Furthermore, 39 K is one of the low- γ quadrupolar nuclides that are extremely difficult to study by NMR at low magnetic fields. Only a few solid-state 39K NMR studies have been reported, and most of those focus on simple inorganic K⁺ salts.¹⁹ Until now, crystallography has been the only biophysical technique capable of directly localizing K⁺ ions bound to biological structures. Here we report the first solid-state ³⁹K NMR detection of K⁺ ions bound to the G-quadruplex structures formed by 5'-tert-butyl-dimethylsilyl-2',3'-O-isopropylidene guanosine (G1), guanosine (G2), and guanosine 5'-monophosphate (G3) (Scheme 1).²⁰ Our strategy for overcoming practical difficulties in studying 39K is to utilize an ultrahigh magnetic field, 19.6 T (830 MHz for ¹H), at the National High Magnetic Field Laboratory (NHMFL).

The lipophilic nucleoside G1 was used as the standard sample for our 39K NMR experiment because G1 self-associates in the presence of K⁺ and Cs⁺ picrate to form a crystallographically defined G-quadruplex.⁷ As shown in Figure 1, the G-quadruplex formed from G1 consists of four G-quartets that are stacked on top of one another to give a structure with a central ion channel. This channel is fully occupied by three collinear K⁺ ions along its central axis. Each of the K⁺ ions is sandwiched by two G-quartets, a structural feature remarkably similar to that recently found in G-rich oligonucleotides.^{8,9} As shown in Figure 2, the ³⁹K magicangle spinning (MAS) spectrum of G1 exhibits a peak centered at -45 ppm. The detailed features in the line shape suggest an overlap of several central-transition powder spectra. Because there are three crystallographically distinct, yet similar, K+ sites inside the G1 quadruplex channel, it is not possible to extract an accurate value

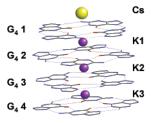


Figure 1. Diagram illustrating the cation binding environment in the G-quadruplex structure formed by G1 self-association.⁷

Scheme 1

for the 39 K nuclear quadrupole coupling constant (C_0) for each of the K⁺ sites. On the basis of the ³⁹K spectra obtained at 19.6 and 11.7 T, we obtained the following estimates: $\delta = -42$ ppm and $C_0 \le 0.7$ MHz. G2 forms a viscous gel in water in the presence of KCl, indicating formation of a highly ordered molecular assembly. An earlier X-ray fiber diffraction study confirmed that the G2 aggregates have a quadruplex structure.²¹ The ³⁹K MAS spectrum of G2 exhibits a peak with $\delta = -45$ ppm and $C_0 \approx 0.7 - 0.8$ MHz. An additional sharp peak at -9 ppm arises from a small excess of KCl. These observations suggest that, in the self-aggregates of G2, the K⁺ ions reside exclusively inside the quadruplex channel in a fashion similar to those in the G1 quadruplex. G3 is among the earliest examples examined by Gellert et al., who first proposed the G-quartet model.²² Subsequent X-ray diffraction studies confirmed that the self-assembly of G3 forms a right-handed quadruplex helix.²³ In contrast to nucleosides G1 and G2, G3 is a mononucleotide where the negatively charged phosphate group is another potential K⁺ binding site. Consequently, in addition to the large KCl signals (-9 ppm and associated spinning sidebands) and the signal attributable to the channel K^+ ions, the ^{39}K MAS spectrum of G3 shows another peak with a line width 3 times greater than that of the signal from the channel K^+ ions. This broad signal (δ \approx -60 ppm) can be assigned to the K⁺ ions bound to the phosphate group. The ratio between the signal areas for the phosphate-bound and channel K⁺ ions is approximately 3:2, much smaller than the 8:1 ratio expected for a G3 quadruplex saturated with K⁺ ions. Our previous study showed that the G-quadruplex channel strongly favors K+ ions, whereas the doubly charged phosphate group of G3 prefers Na⁺ ions over K⁺ ions.¹³ As the G-quadruplex sample containing G3 was prepared in the presence of both K⁺ and Na⁺

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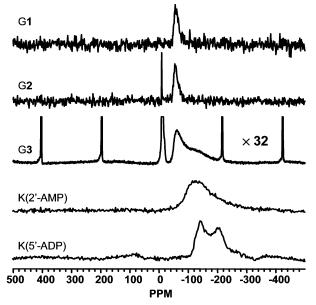


Figure 2. Experimental ³⁹K MAS NMR spectra at 19.6 T. All solid-state NMR experiments were performed with a narrow-bore magnet (31 mm) and a Bruker Avance console at NHMFL operating at 38.72 MHz for ³⁹K nuclei. A home-built MAS probe equipped with a 4-mm stator was used. Reported ³⁹K chemical shifts were referenced to KBr(s), $\delta = 0$ ppm. Other experimental details: G1, 10-kHz spinning, 7500 transients, 2-s recycle delay; G2, 10-kHz spinning, 12 864 transients, 0.5-s recycle delay; G3, 8-kHz spinning, 100 000 transients, 1-s recycle delay; K(2'-AMP), 8-kHz spinning, 34 956 transients, 2-s recycle delay; K(5'-ADP), 8-kHz spinning, 45 364 transients, 2-s recycle delay.

ions,²⁰ the central channel is clearly filled with K⁺ ions; however, there are likely to be a considerable amount of Na⁺ ions that remain bound to the phosphate groups. To further confirm this spectral assignment, we obtained ³⁹K MAS spectra for hydrated K salts of adenosine 2'-monophosphate and adenosine 5'-diphosphate, K(2'-AMP)·1.5H₂O and K(5'-ADP)·2H₂O. The K⁺ ion in K(2'-AMP) is coordinated to two phosphate oxygens, two water molecules, and two hydroxyl groups from the ribose. 24 The 39 K spectrum of K(2'-AMP) exhibits a broad peak with $\delta = -55$ ppm and $C_0 = 1.85$ MHz, in excellent agreement with that observed for the phosphatebound K^+ ions in G3. The K^+ ion in K(5'-ADP) is coordinated to seven neighbors: four phosphate oxygens, one water, one hydroxyl, and an N atom from the adenine base.²⁵ The ³⁹K MAS spectrum of K(5'-ADP) exhibits a clear second-order quadrupole line shape, with $\delta = -105$ ppm and $C_0 = 2.05$ MHz. These parameters are quite different from those observed for the K+ ions in Gquadruplexes, reflecting the unusual K^+ coordination in K(5'-ADP).

In summary, we have obtained unambiguous 39 K NMR signatures for the K^+ ions bound to G-quadruplex structures. This is an important first step toward solid-state 39 K NMR studies of telomeric DNA. The rich spectral information in the 39 K NMR spectra of K-nucleotide systems suggests that solid-state NMR is a viable new method for detecting K^+ ions in biomolecular systems.

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