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## NMR Detection of Protein <sup>15</sup>N Spins near Paramagnetic Lanthanide Ions

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The strong paramagnetism associated with lanthanide ( $Ln^{3+}$ ) ions provides a unique source of structural information available for proteins that either have a natural lanthanide-binding site<sup>1,2</sup> or are proteins site specifically tagged with an engineered lanthanidebinding group.<sup>3,4</sup> It allows rapid 3D structure determinations of protein—protein and protein—ligand complexes,<sup>2,5</sup> detailed structural studies near the lanthanide binding site,<sup>6</sup> and the measurement of residual dipolar couplings.<sup>7</sup> Among the paramagnetically induced NMR effects, pseudocontact shifts (PCS) of amide groups contain particularly useful structural information, because they report on the location of the nuclear (i.e., <sup>15</sup>N and <sup>1</sup>H<sup>N</sup>) spins with respect to the metal's magnetic susceptibility anisotropy ( $\Delta \chi$ ) tensor anchored in the molecular frame:<sup>8</sup>

$$\Delta \delta^{\text{PCS}} = \frac{1}{12\pi r^3} \Big[ \Delta \chi_{\text{ax}} (3\cos^2\theta - 1) + \frac{3}{2} \Delta \chi_{\text{rh}} \sin^2\theta \cos(2\phi) \Big]$$
(1)

where  $\Delta \chi_{ax}$  and  $\Delta \chi_{rh}$  are the axial and rhombic components of the  $\Delta \chi$  tensor, and  $(r,\varphi,\theta)$  are the spherical coordinates of the nuclear spin in the tensor's principal axis system. Lanthanides like Tb<sup>3+</sup> or Dy<sup>3+</sup> generate large PCS that are measurable for nuclear spins as far as 40 Å from the metal ion.<sup>9</sup>

As a drawback, the strongest paramagnetic  $Ln^{3+}$  ions are invariably associated with pronounced paramagnetic relaxation enhancements (PRE). In particular, transverse PRE of <sup>1</sup>H spins render amide groups within about 15 Å of a Dy<sup>3+</sup> ion and, hence, the entire metal binding site unobservable in <sup>15</sup>N-HSQC spectra. For most lanthanides, transverse PRE are dominated by the Curie mechanism<sup>8</sup> and, therefore, can be approximated by

$$\Delta R_2^{\rm PRE} = \frac{1}{5\pi r^6} \chi_{\rm iso}^2 \gamma^2 B_0^2 \tau_{\rm R}$$
 (2)

where  $\chi_{iso}$  is the metal's isotropic magnetic susceptibility,  $\gamma$  is the gyromagnetic ratio of the nuclear spin,  $B_0$  is the magnetic field strength, and  $\tau_R$  is the molecular rotational correlation time. Because of the strong dependence of PRE on the metal-nuclear distance r, additional spectra with much weaker paramagnetic ions such as Ce<sup>3+</sup> have to be acquired to yield PCS information close to the metal center.

Alternatively, the scaling of PRE with  $\gamma^2$  has been exploited in a range of protonless <sup>13</sup>C-detected experiments.<sup>10 13</sup>C spins are about 16-fold less sensitive to PRE effects than <sup>1</sup>H spins, so that the same PRE prevails 1.6-fold closer to the metal ion. Correspondingly, <sup>15</sup>N spins would be observable more than 2-fold closer.<sup>6,11</sup> Much of this gain is, however, lost by the intrinsically lower sensitivity of these heteronuclear spins and the difficulty to assign broad resonances by correlation spectra with fairly long magnetization transfer periods.<sup>6</sup>

Here we present an approach that combines the advantage of <sup>1</sup>H detection with the favorable relaxation properties of <sup>15</sup>N and the resolution of the <sup>15</sup>N-HSQC spectrum to measure PCS for <sup>15</sup>N spins



**Figure 1.** Out-and-back N<sub>z</sub>-exchange experiment. Narrow and wide bars represent rf pulses with flip angles of 90° and 180°, respectively, applied with phase *x* unless otherwise indicated. The cycled phases were  $\phi_1(y, -y)$ ,  $\phi_2(x)$ ,  $\phi_3(2x, 2(-x))$ ,  $\phi_4(y)$ , and  $\phi_{rec}(x, 2(-x), x)$ , and quadrature detection was achieved with States-TPPI applied to  $\phi_2$  and  $\phi_4$ . The delays  $\tau_{\rm H}$ ,  $\tau_{\rm N}$ , and  $\delta$  were set to 2.5, 2.75, and 1.5 ms, respectively. N was set to 160 using a MLEV-16 cycle for the <sup>1</sup>H pulses, resulting in mixing times ( $\tau_{\rm m1} = \tau_{\rm m2}$ ) of 486 ms. The pulsed field gradients were applied along *z* with strengths ( $G_i$ ) of 23.2, 8.7, 11.6, 14.5, 20.3, and 11.6 G/cm, and water suppression was achieved using the 3-9-19 sequence. Although the scheme was formally designed for water flip-back, the water resonance becomes saturated for long mixing periods. Omission of the <sup>1</sup>H pulses during  $\tau_{\rm m}$  resulted in a 20% increase of  $R_1$  relaxation rates.

as close as 6 Å from the Dy<sup>3+</sup> ion in a 30 kDa protein complex. The experiment relies on the chemical exchange of metal ions in a sample prepared with a mixture of diamagnetic and paramagnetic metals, and is implemented as an out-and-back experiment, where magnetization is stored as PRE-insensitive, longitudinal <sup>15</sup>N magnetization during two mixing times (Figure 1).

The use of two rather than one  $N_z$  mixing period distinguishes the experiment from conventional heteronuclear ( $N_z$ ) exchange spectroscopy.<sup>12,13</sup> This allows the generation of cross-peaks by a pathway (see Supporting Information) where magnetization is transferred from the diamagnetic to the paramagnetic state during  $\tau_{m1}$ , frequency-labeled during  $t_1$ , and transferred back to the diamagnetic state during  $\tau_{m2}$ , avoiding the presence of transverse magnetization in the paramagnetic state at any other time than  $t_1$ . Consequently, the observability of <sup>15</sup>N spins is limited by <sup>15</sup>N PRE rather than <sup>1</sup>H PRE.

The fraction of  $N_z$  magnetization transferred to the paramagnetic state during  $\tau_{m1}$  and back during  $\tau_{m2}$  is given by

$$M(\tau_{\rm m})/M_{\rm tot} = f_{\rm d}^2 f_{\rm p} (1 - e^{-k_{\rm ex}\tau_{\rm m}})^2 e^{-2R_{\rm l}\tau_{\rm m}}$$
(3)

where  $M_{\rm tot}$  denotes the total N<sub>z</sub> magnetization at the start of  $\tau_{\rm m1}$ ,  $f_{\rm d}$  and  $f_{\rm p}$  are the molar fractions of diamagnetic and paramagnetic protein, respectively,  $k_{\rm ex}$  is the rate of metal exchange,  $R_1$  is the longitudinal <sup>15</sup>N relaxation rate (assumed to be the same in both states), and it is assumed that  $\tau_{\rm m} = \tau_{\rm m1} = \tau_{\rm m2}$ . To minimize losses, the experiment requires that  $k_{\rm ex}$  is not much slower than  $R_1$ .

We applied the experiment to the N-terminal domain of the exonuclease subunit  $\epsilon$  of *E. coli* DNA polymerase III ( $\epsilon$ 186) in complex with the subunit  $\theta$ . The active site of  $\epsilon$ 186 binds two divalent ions<sup>14</sup> that can be replaced by a single Ln<sup>3+</sup> ion.<sup>15</sup> Although Ln<sup>3+</sup> binding is quite tight with dissociation constants in the low  $\mu$ M range ( $k_{ex} < 1 \text{ s}^{-1}$ ), catalysis of metal exchange by excess metal<sup>15</sup> allows tuning of  $k_{ex}$  to values that compete with  $R_1$  while



Figure 2. Selected strips from the 3D out-and-back Nz-exchange spectrum of the 30 kDa  $\epsilon$ 186/ $\theta$  complex, containing <sup>2</sup>H/<sup>15</sup>N-labeled  $\epsilon$ 186. Parameters used: 25 °C, 0.8 mM  $\epsilon$ 186/ $\theta$ , equimolar ratio of Dy<sup>3+</sup> and La<sup>3+</sup>, 20 mM Tris (pH 7.2), 100 mM NaCl, Bruker Avance 800 MHz NMR spectrometer with TCI cryoprobe,  $\tau_{m1} = \tau_{m2} = 486$  ms,  $96 \times 64 \times 1280$  complex points, spectral widths of 32 ppm (<sup>15</sup>N) and 20 ppm (<sup>1</sup>H), total experiment time 40 h. The strips are centered on the  ${}^{15}N(F_2)$  and  ${}^{1}H^N(F_3)$  frequencies of the diamagnetic peaks of residues 169-176. Lines connect diagonal and exchange peaks and are labeled with the chemical shift difference which corresponds to the <sup>15</sup>N PCS.

avoiding excessive line-broadening because of the exchange or unspecific binding of lanthanide ions. By stepwise addition of a 1:1 mixture of Dy<sup>3+</sup> and La<sup>3+</sup> to  $\epsilon$ 186/ $\theta$  we adjusted  $k_{ex}$  to about 5 s<sup>-1</sup>, that is, 10-fold larger than  $R_1$ . Under these conditions, about 35% of initial diamagnetic N<sub>z</sub> magnetization can be transferred to the paramagnetic state (and vice versa) during a single mixing period of 0.5 s, whereas 40% are retained in the diamagnetic state, and 25% are lost owing to  $R_1$  relaxation. This presents a substantial gain over an experiment that starts from <sup>15</sup>N polarization (see Supporting Information).

Figure 2 shows strips from the 3D out-and-back N<sub>z</sub>-exchange spectrum for the helical segment from Q169 to L176. Each strip contains diagonal ( $F_1 = F_2$ ) and exchange peaks (where  $F_1$  reports the <sup>15</sup>N frequency of the paramagnetic state). The frequency difference between both peaks corresponds to the <sup>15</sup>N PCS. PCS values of up to 40 ppm and 15N line widths of up to 130 Hz were measured for the  $\epsilon 186/\theta$ /Dy<sup>3+</sup> complex for a total of 156 residues of  $\epsilon$ 186. Notably, 61 of these could not be observed in the <sup>15</sup>N-HSQC spectrum of the pure  $\epsilon 186/\theta$ /Dy<sup>3+</sup> complex owing to excessive <sup>1</sup>H PRE. For all remaining residues of  $\epsilon$ 186, for which the resonance assignments are known in the diamagnetic  $\epsilon 186/\theta/$ La<sup>3+</sup> complex, the <sup>15</sup>N-Ln<sup>3+</sup> distances are shorter than 6 Å, diagonal and exchange peaks are overlapped, or the peaks are already weak in the diamagnetic <sup>15</sup>N-HSQC spectrum.

The spectrum also contains the expected diagonal and exchange peaks that are detected on the paramagnetic <sup>1</sup>H<sup>N</sup> resonance for residues with r > 15 Å. In the projection along  $F_2$ , the four peaks form a rectangle and are therefore easy to identify.

For many of the residues close to the metal ion, the <sup>15</sup>N PCS values measured in the  $\epsilon 186/\theta/Dy^{3+}$  complex were found to deviate substantially from values back-calculated using previously determined  $\Delta \chi$  tensors of  $\epsilon 186/\theta$ /Dy<sup>3+</sup>.<sup>16</sup> This may be due to chemical exchange between multiple Ln<sup>3+</sup> binding sites (up to three different, closely spaced  $Ln^{3+}$  sites were observed in crystals of  $\epsilon 186$  soaked with  $Dy^{3+}$ , ref 17) or due to structural differences between  $\epsilon$ 186 in the single crystal<sup>14</sup> and the  $\epsilon 186/\theta$  complex in solution. Notably, corresponding deviations in <sup>15</sup>N PCS values were also observed for the  $\epsilon 186/\theta/\text{Ce}^{3+}$  complex. This indicates that the deviations report on structural rather than electronic features (see Supporting Information).

Ln<sup>3+</sup> exchange rates on the time scale of the out-and-back N<sub>2</sub>exchange experiment have also been observed for lanthanidebinding peptide tags<sup>4</sup> (Su, X.C., personal communication). This makes the new experiment widely applicable for the assignment of <sup>15</sup>N resonances and measurement of <sup>15</sup>N PCS of proteins labeled with paramagnetic lanthanides.

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Supporting Information Available: Scheme of magnetization transfer pathways, comparison with an experiment starting from <sup>15</sup>N magnetization, correlation of deviations between experimental and backcalculated <sup>15</sup>N PCS for Ce<sup>3+</sup> versus Dy<sup>3+</sup>,  $\Delta \chi$  tensor parameters, and a table of  ${}^{1}\text{H}{}^{N}$  and  ${}^{15}\text{N}$  chemical shifts of  $\epsilon 186$  in complex with  $\theta$  and  $La^{3+}$ ,  $Dy^{3+}$  or  $Ce^{3+}$ . This material is available free of charge via the Internet at http://pubs.acs.org.

## References

- (a) Bentrop, D.; Bertini, I.; Cremonini, M. A.; Forsen, S.; Luchinat, C.; Malmendal, A. *Biochemistry* 1997, *36*, 11605–11618. (b) Biekovsky, R. R.; Muskett, F. W.; Schmidt, J. M.; Martin, S. R.; Browne, J. P.; Bayley, P. M.; Feeney, J. *FEBS Lett.* **1999**, *460*, 519–526. (c) Contreras, M. A.; Ubach, J.; Millet, O.; Rizo, R.; Pons, M. J. Am. Chem. Soc. 1999, 121, 8947-8948. (d) Ma, C.; Opella, S. J. J. Magn. Reson. 2000, 146, 381-384
- Nitz, M.; Imperiali, B.; Schwalbe, H. J. Am. Chem. Soc. 2003, 125, 13338–13339. (c) Ikegami, T.; Verdier, L.; Sakhaii, P.; Grimme, S.; **2004**, *29*, 339–349. (d) Pintacuda, G.; Moshref, A.; Leonchiks, A.; Sharipo, A.; Otting, G. J. Biomol. NMR 2004, 29, 351-361. (e) Prudencio, M.; Rohovec, J.; Peters, J. A.; Tocheva, E.; Boulanger, M. J.; Murphy, M. E. P.; Hupkes, H.-J.; Kosters, W.; Impagliazzo, A.; Ubbink, M. *Chem.–Eur. J.* **2004**, *10*, 3252–3260.
- (4) Su, X. C.; Huber, T.; Dixon, N. E.; Otting, G. ChemBioChem 2006, 7, 1599 - 1604
- (5) John, M.; Pintacuda, G.; Park, A. Y.; Dixon, N. E.; Otting, G. J. Am. Chem. Soc. 2006, 128, 12910–12916.
- (6) Balayssac, S.; Jiminez, B.; Piccioli, M. J. Biomol. NMR 2006, 34, 63-
- (7) Tolman, J. R.; Flanagan, J.; Kennedy, M. A.; Prestegard, J. H. Proc. Natl. Acad. Sci. U.S.A. 1995, 92, 9279–9283.
- (8) Bertini, L.; Luchinat, C.; Parigi, G. Prog. NMR Spectrosc. 2002, 40, 249-273
- (9) Allegrozzi, M.; Bertini, I.; Janik, M. B. L.; Lee, Y.-M.; Liu, G.; Luchinat, C. J. Am. Chem. Soc. 2000, 122, 4154–4161.
  (10) Bermel, W.; Bertini, I.; Felli, I. C.; Piccioli, M.; Pierattelli, R. Prog. NMR
- Spectrosc. 2006, 48, 25-45 (a) Sadek, M.; Brownlee, R. T. C. J. Magn. Reson., Ser. B 1995, 109, 70–75. (b) Xia, B.; Wilkens, S. J.; Westler, W. M.; Markley, J. L. J. Am. (11)
- Chem. Soc. 1998, 120, 4893-4894. (12) Farrow, N. A.; Zhang, O.; Forman-Kay, J.; Kay, L. J. Biomol. NMR 1994,
- 4.727 (13) John, M.; Headlam, M.; Dixon, N. E.; Otting, G. J. Biomol. NMR, in
- ress. (14) Hamdan, S.; Carr, P. D.; Brown, S. E.; Ollis, D. L.; Dixon, N. E. Structure, 2002, 10, 535–546.
- (15) Pintacuda, G.; Keniry, M. A.; Huber, T.; Park, A. Y.; Dixon, N. E.; Otting,
- G. J. Am. Chem. Soc. 2004, 126, 2963–2970.
   Schmitz, C.; John, M.; Park, A. Y.; Dixon, N. E.; Otting, G.; Pintacuda, G.; Huber, T. J. Biomol. NMR 2006, 35, 79-87.
- (17) Park, A. Y. Ph.D. Thesis. Australian National University, Australia, 2006.

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