

Cytosine–guanine base pairing in a hydrogen-bonded complex of stable open-shell molecules with $S = 1$ spins†

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A bio-inspired motif of hydrogen-bonded nucleobases is introduced to stable organic biradical crystals. We have synthesized a hydrogen-bonded complex based on both a cytosine-substituted nitronyl nitroxide biradical (**1**) and a closed-shell alkyl-substituted guanine (**2**). From single-crystal X-ray structure analyses, the cytosine and guanine moieties are found to form a Watson–Crick pair with threefold hydrogen bonds. Magnetic susceptibility measurements reveal that the cytosine-substituted biradical **1** has a triplet ($S = 1$) ground state with a singlet–triplet energy gap of $2J/k_B = 27.0$ K. This complex is the first example of Watson–Crick type base pair possessing stable high-spin organic radical moieties, which is fully characterized by single-crystal X-ray structure analysis. A face-to-face overlapping is found between the planar nucleobases, while side-by-side hydrogen bonds are found between the Watson–Crick pairs. Thus, the introduction of the radical substituent into the nucleobase results in little disturbance to the molecular arrangement usually found in nucleobase molecules: the ground-state spin multiplicity of biradical **1** is maintained.

Introduction

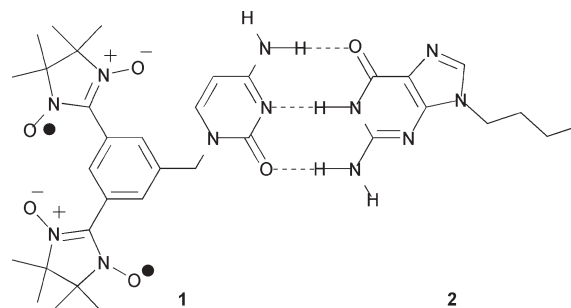
The last few decades have witnessed rapid development of molecule-based magnetism.¹ Molecular arrangement in a crystalline solid is closely related to bulk magnetic properties of molecular assemblages such as ferromagnetism underlain by intermolecular magnetic interactions. Thus, an important issue in molecule-based magnetism is control of molecular packing, or relative arrangement of open-shell building block molecules in a crystalline solid state. Molecular packing can be partially controlled by giving the building block molecules specific noncovalent bondings. Among intermolecular noncovalent bondings, hydrogen bonding is promising in controlling the molecular arrangement in the crystalline solid. Hydrogen-donating and accepting molecules such as phenol, carboxylic acid, and pyridine derivatives have been introduced to the stable radical family of nitroxide and imino or nitronyl nitroxide.² Nucleobases (nucleic acid base) such as guanine, cytosine, adenine and thymine found in a DNA duplex can be further fascinating carriers of noncovalent bondings as they can form complementary hydrogen bonds with high selectivity and directionality, as compared with other hydrogen-donating and accepting molecular systems. Several examples have been found in which stable radicals of nitroxide are introduced to nucleobases in crystalline solid states.^{3–6} Solution chemistry and ESR spectroscopy have been reported for spin-labeled

nucleosides and nucleotides.^{7,8} In this study, we have synthesized an open-shell substituted heteromolecular complex of cytosine (**1**) and guanine (**2**) derivatives (Scheme 1). The cytosine is substituted with a stable nitronyl nitroxide biradical with a spin $S = 1$ in the ground state. This is the first example of Watson–Crick type base pair possessing stable high-spin ($S > 1/2$) organic radical moieties,⁹ which is fully characterized by single-crystal X-ray structure analysis and bulk magnetic susceptibility. The influence of the radical substituent on the molecular arrangement by the hydrogen bonding of the nucleobase moieties is examined.

Experimental

Synthesis

The cytosine-substituted biradical **1** was synthesized by a direct coupling of cytosine with a 5-chloromethylphenyl derivative of nitronyl nitroxide **3**,¹⁰ as shown in Scheme 2. The precursor **3** has been prepared from hexatriene using literature procedures.^{11,12} The precursor **3** (100 mg, 0.23 mmol) was added to a suspension of cytosine (38.8 mg, 0.35 mmol), K_2CO_3 (46.8 mg,

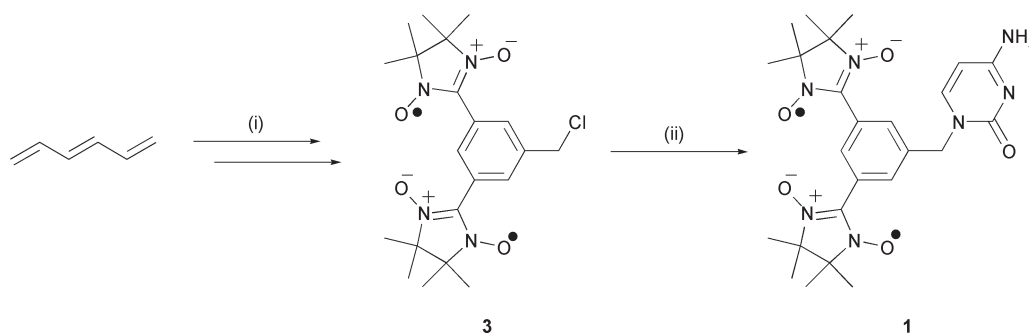


Scheme 1 Nucleobase pair substituted with nitronyl nitroxide.

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† Electronic supplementary information (ESI) available: ORTEP drawings and results of DFT molecular orbital calculations. See DOI: 10.1039/b706521n



Scheme 2 Synthesis of **1**. Steps: (i) procedures reported in ref. 10; (ii) cytosine, K_2CO_3 and KI in acetonitrile for 18 h.

0.34 mmol), and KI (53.6 mg, 0.32 mmol) in dry acetonitrile (6 ml). The reaction mixture was stirred at 80 °C for 18 h. After filtration, the reaction solution was concentrated and chromatographed on silica gel using acetone–methanol (7 : 1 in volume) as an eluent to yield a blue solid of **1** (66.3 mg, 58.2%). Elemental analysis found: C 57.52, H 6.57, N 17.60. Calcd. for $C_{25}H_{33}N_7O_5 + CH_3OH$: C 57.44, H 6.86, N 18.04%.

The closed-shell guanine derivative **2** was synthesized by the literature method.¹³ Single crystals of the complex of **1** and **2** were obtained by recrystallization from a mixed solution of ethanol, chloroform and diethyl ether. Elemental analysis found: C 55.58, H 6.72, N 21.61. Calcd. for $C_{34}H_{46}N_{12}O_6 + C_2H_5OH + H_2O$: C 55.23, H 6.95, N 21.47%.

ESI-MS

Electrospray ionization mass spectrometry (ESI-MS) experiments were carried out on a JEOL JMS700T Tandem Mstation. An equal-volume mixture of methanol and chloroform was used as the solvent for electrospray. The spray voltage was 2.0 kV in the positive-ion mode.

Crystal structure analysis

Single crystal diffraction data of the complex of **1** and **2** were collected on a Rigaku Mercury CCD diffractometer with graphite monochromated Mo $K\alpha$ radiation at 193 K. The crystal structure was solved by using direct methods (SHELXS97)¹⁴ in a program package *CrystalStructure*.¹⁵ Anisotropic thermal parameters analysis was applied for non-hydrogen atoms. Hydrogen atoms with isotropic thermal parameters were refined as riding models. From powder X-ray diffraction experiments, we have confirmed that the crystal structure of the polycrystalline sample used in magnetic susceptibility measurements is the same as that determined in the single crystal X-ray experiments.

Magnetic susceptibility measurements

The static paramagnetic susceptibility χ_p was measured for polycrystals on a SQUID magnetometer with an applied magnetic field of 0.1 T in the temperature range of 1.9–300 K. The magnetic responses were corrected with diamagnetic blank data of the sample holder obtained separately. The diamagnetic contribution of the sample itself was estimated from Pascal's constants.

Results and discussion

ESI-MS

Formation of a complex of **1** and **2** in solution was examined by ESI-MS. The positive ion mode spectrum of **1** + **2** in a 1 : 1 ratio in a mixed solution of methanol and chloroform (1 : 1) showed three doublet peaks at $m/z = (208, 230)$, (512, 534), and (720, 742), as shown in Fig. 1. The doublets of $m/z = (208, 230)$ and (512, 534) come from $\{(2 + H)^+, (2 + Na)^+\}$ and $\{(1 + H)^+, (1 + Na)^+\}$, respectively. The largest in intensity of the three doublets is $m/z = (720, 742)$, as assigned to a one-to-one complex of **1** and **2** $\{(1 + 2 + H)^+, (1 + 2 + Na)^+\}$. This result is indicative of the formation of a stable 1 : 1 complex of **1** and **2** in solution. An additional peak is found at m/z 851, which is assignable to a tetramer of **2**, $(2_4 + Na)^+$.^{16,17}

Crystal structure

The crystalline solid of the complex of **1** and **2** was stable under aerated conditions at room temperature. The crystallographic data for the complex of **1** and **2** are listed in Table 1. The crystal contains one mole of ethanol as crystal solvent.¹⁸ Fig. 2 depicts the asymmetric unit of the hydrogen-bonded complex of **1** : **2** in a 1 : 1 ratio with the atom numbering. CCDC reference numbers 609687. For crystallographic data in CIF or other electronic format see DOI: 10.1039/b706521n

The cytosine and guanine moieties are found to form a Watson–Crick pair with threefold hydrogen bonds, (i) N7H–O6 = 2.867(5) Å, (ii) N6–HN8 = 2.947(5) Å and (iii) O5–HN9 = 2.942(4) Å. The hydrogen bond lengths fall within

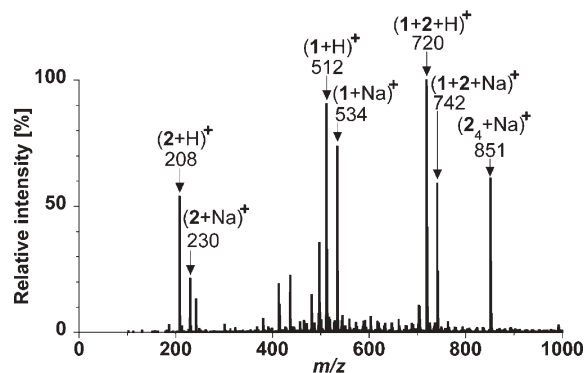


Fig. 1 Positive-ion ESI-MS of **1** + **2** in a 1 : 1 ratio in a mixed solution of methanol and chloroform (1 : 1).

Table 1 Crystallographic data for hydrogen-bonded complex (**1** + **2**)

Formula	C ₃₆ H ₅₂ N ₁₂ O ₇ (1 + 2 + EtOH)
Crystal system	Triclinic
Space group	<i>P</i> -1
<i>a</i> /Å	7.596(6)
<i>b</i> /Å	12.988(1)
<i>c</i> /Å	20.897(2)
α /°	100.891(1)
β /°	95.406(1)
γ /°	95.522(1)
<i>V</i> /Å ³	2002(3)
<i>Z</i>	2
ρ_{calc} /g cm ⁻³	1.269
<i>T</i> /K	143
Unique reflections used (All reflections)	7900
<i>R</i> (<i>I</i> > 2 σ (<i>I</i>)) ^a	0.0999
<i>R</i> _{w2} (All reflections) ^b	0.2666
Goodness of fit	1.001

^a $R = \frac{\sum ||F_o| - |F_c||}{\sum |F_o|}$.

^b $R_{w2} = \left[\frac{\sum w(F_o^2 - F_c^2)^2}{\sum w(F_o^2)} \right]^{\frac{1}{2}}$,
 $w = 1/[0.0022F_o^2 + 1.0000(F_o^2)]/(4F_o^2)$

the typical values of Watson–Crick pairs of cytosine–guanine complexes, which were studied previously by X-ray single-crystal structure analyses.^{19,20}

The Watson–Crick pairs are hydrogen-bonded to form a ribbon-shaped structure ((iv) N7H–N11 = 3.006(6) Å, (v) O5–HN9 = 2.905(5) Å) within the molecular plane of the hydrogen-bonded nucleobase moieties, as depicted in Fig. 3(a). The side-by-side hydrogen-bonding gives an alternating arrangement of the cytosine and guanine moieties (–C–G–C–G–) along the *a* + *b* axis. This type of side-by-side hydrogen-bonding mode has been found in nonpolymeric cytosine–guanine complexes.¹⁹ In oligonucleotide duplexes, hydrogen-bonded nucleobase moieties are stacked with the molecular planes facing each other.²⁰ The face-to-face overlapping of planar nucleobases is found in the neighboring cytosine and guanine moieties of the complex as well, resulting in the columnar structure in the direction nearly perpendicular to the hydrogen-bonding molecular plane, as shown in

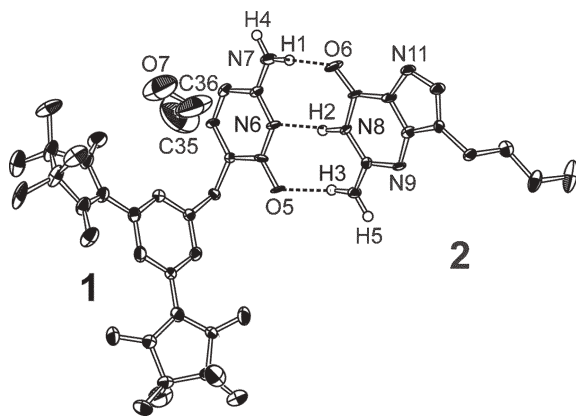


Fig. 2 ORTEP²⁵ view of the Watson–Crick pair (cytosine–guanine pair) with the thermal ellipsoids at the 50% probability level. The hydrogen atoms are omitted for clarity except those in the hydrogen bonding moiety as shown by the dotted lines.

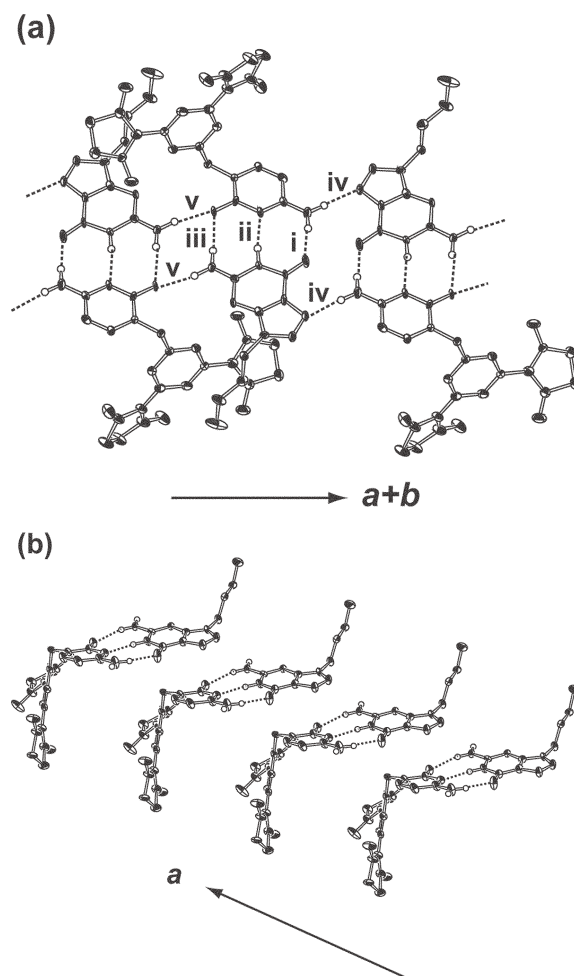


Fig. 3 (a) Hydrogen-bonded network along the *a* + *b* axis. The intermolecular hydrogen bonds are labeled i through v. (b) Stacking of the complex running along the *a* axis. The methyl groups are omitted for clarity. The thermal ellipsoids are given at the 50% probability level.

Fig. 3(b).²¹ Thus, introduction of the radical substituent into the nucleobase results in little disturbance to the molecular arrangement usually found in nucleobase molecules or intermolecular hydrogen bonding motifs other than the Watson–Crick pairing.

The phenyl and the cytosine rings in **1** have a large dihedral angle of 100.8° around the methylene bridge. It is expected that the delocalization of unpaired electron spin in **1** is truncated at the methylene bridge. It was found from molecular orbital calculations that the cytosine moiety has little spin density.²² Thus, the cytosine group in **1** should play a role primarily in determining the molecular packing structure instead of propagating intermolecular magnetic interactions.

Magnetic properties

The temperature dependence of paramagnetic susceptibility χ_p measured for randomly oriented polycrystals of the complex is shown in Fig. 4 in the $\chi_p T$ versus *T* plots.²³ The $\chi_p T$ value of 0.75 emu K mol⁻¹ at 300 K corresponds to 2 moles of free

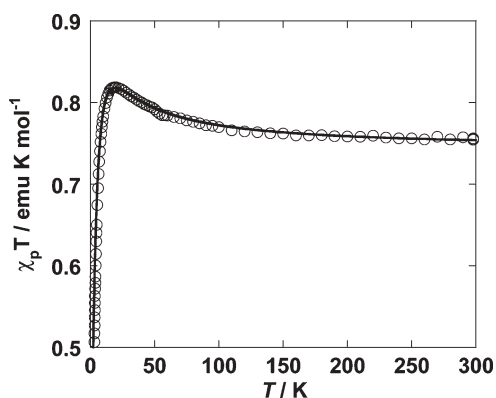


Fig. 4 Temperature dependence of paramagnetic susceptibility χ_p of the complex in the $\chi_p T$ versus T plots. The solid line is calculated from eqn (1).

$S = 1/2$ spins with the g -factor of 2.0, indicating that the purity of the polycrystals is satisfactory. The $\chi_p T$ value increases gradually as the temperature is lowered, and the value decreases rapidly below 20 K, indicating that both a strong ferromagnetic interaction and a weak antiferromagnetic one undergo in the crystal. The ferromagnetic interaction is naturally ascribed to the intramolecular exchange interaction in the biradical **1**, as found in precursor **3** with a similar structure,¹⁰ while the weak antiferromagnetic interaction is attributed to intermolecular contacts. An estimate of the intra- and inter-molecular exchange interactions is acquired by assuming the singlet–triplet equilibrium,²⁴ given as

$$\chi_p = \frac{\alpha N_A g^2 \mu_B^2}{k_B(T-\theta)} \times \frac{2}{3 + \exp(-2J/k_B T)} \quad (1)$$

where $2J$ denotes the intramolecular exchange interaction, *i.e.*, the singlet–triplet energy gap of **1**. The intermolecular antiferromagnetic interaction is approximated by a mean-field signified by the Weiss constant θ . N_A , k_B , and α in eqn (1) are the Avogadro constant, Boltzmann constant and purity, respectively. The observed $\chi_p T$ is well reproduced by assuming $2J/k_B = 27.0 \pm 0.3$ K, $\theta = -2.3 \pm 0.1$ K, $g = 2.006$, and $\alpha = 0.99$. An intermolecular contact around the nitroxide oxygen atom with a large spin density is found at O2–C18 = 3.158(8) Å between two neighboring inversion-related molecules **1**, which is one of the possible contacts contributing to the intermolecular antiferromagnetic interactions of $\theta = -2.3$ K. The positive J value shows that the ground state of the biradical **1** is triplet ($S = 1$) in the complex.

Conclusions

We have synthesized the first example of a cytosine–guanine complex of the Watson–Crick type, which carries a stable biradical in the triplet ($S = 1$) ground state. The X-ray crystallographic analysis has shown that the complex possesses side-by-side hydrogen bonding and face-to-face stacking extending over the Watson–Crick pairs in the crystal. Introduction of the large radical substituent into the nucleobase results in little disturbance to the intermolecular hydrogen bonding motifs usually found in cytosine–guanine pairings in

crystalline solid states. Thus, biradical **1** maintains the ground-state triplet in the crystal.

Acknowledgements

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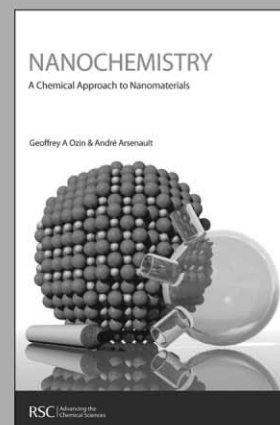
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