

## FT-IR and $^1\text{H}$ NMR Spectroscopic Evidence of Sugar Ring Conformational Change in $\text{NH}_4\text{GpG}$ on Complexation to form $\text{cis-}[\text{Pt}(\text{NH}_3)_2(\text{GpG})]^+$

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### Abstract

The conformational change of the ribose ring in  $\text{NH}_4\text{GpG}$  and  $\text{cis-}[\text{Pt}(\text{NH}_3)_2(\text{GpG})]^+$  was confirmed by FT-IR spectroscopic evidence as being  $\text{C}2'$ -endo,  $\text{C}3'$ -endo, anti, gg sugar ring pucker in the solid state. These results were compared with  $^1\text{H}$  NMR spectral data in aqueous solution. The FT-IR spectrum of  $\text{NH}_4\text{GpG}$  shows marker bands at  $802\text{ cm}^{-1}$  and  $797\text{ cm}^{-1}$  which are assigned to the  $\text{C}3'$ -endo, anti, gg sugar-phosphate vibrations of ribose ( $-\text{pG}$ ) and ribose ( $\text{Gp}-$ ), respectively. The FT-IR spectrum of  $\text{cis-}[\text{Pt}(\text{NH}_3)_2(\text{GpG})]^+$  (with  $\text{N}7-\text{N}7$  chelation in the  $\text{GpG}$  sequence) shows a marker band at  $800\text{ cm}^{-1}$  which is assigned to the  $\text{C}3'$ -endo, and a new shoulder band at  $820\text{ cm}^{-1}$  related to a  $\text{C}2'$ -endo ring pucker. The ribose conformation of ( $-\text{pG}$ ) moiety in  $\text{NH}_4\text{GpG}$ ,  $\text{C}3'$ -endo, anti, gg changes into  $\text{C}2'$ -endo, anti, gg when a platinum atom is chelated to  $\text{N}7-\text{N}7$  in the  $\text{GpG}$  sequence.

### Introduction

Until now, only three oligonucleotide–platinum complexes containing the complex  $\text{d}(-\text{GpG}-)\cdot\text{cis-Pt}$  have been the subject of conformational analysis in aqueous solution [1]. In all adducts the  $\text{d}(-\text{GpG}-)\cdot\text{cis-Pt}$  fragments appear to adopt a geometry in which the bases are anti, gg with respect to the sugar rings. The 5'-terminal ( $\text{Gp}-$ ) ring adopts a 100%  $\text{C}3'$ -endo conformation, whereas the 3'-terminal ( $-\text{pG}$ ) ring adopts about 70–90%  $\text{C}2'$ -endo conformation. In the solid state, FT-IR spectroscopy has also been useful to identify the sugar conformational change as previously reported [2].

In the present work, the conformational change of the ribose ring in  $\text{NH}_4\text{GpG}$  and  $\text{cis-}[\text{Pt}(\text{NH}_3)_2(\text{GpG})]^+$  was confirmed by FT-IR spectroscopic evidence being  $\text{C}2'$ -endo,  $\text{C}3'$ -endo, anti, gg sugar ring pucker in the solid state. These results were compared with  $^1\text{H}$  NMR (400 MHz) data in aqueous

solution. The FT-IR spectra were compared to those previously reported [2], and all the platinum complexes were prepared according to the methods described in the literature [3] except that the isolation of the complex,  $\text{cis-}[\text{Pt}(\text{NH}_3)_2(\text{GpG})]^+$  was obtained from the crude reaction mixture which was evaporated at  $40^\circ\text{C}$  to reduce the volume and then precipitated with acetone.

### Results and Discussion

The ribose ring pucker in RNA double helix is normally  $\text{C}3'$ -endo, anti, gg [4] as is found by X-ray analyses in  $\text{NaGpC}$  and  $\text{CaGpC}$  [5]. These results strongly suggest that the two ribose rings in  $\text{NH}_4\text{GpG}$  adopt a  $\text{C}3'$ -endo conformation, as is also found by FT-IR spectroscopy which also supports the  $\text{C}3'$ -endo conformation in both ( $\text{Gp}-$ ) and ( $-\text{pG}$ ) ribose rings as is shown in Fig. 1.

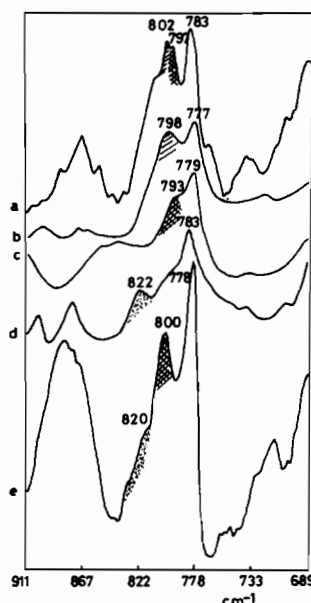


Fig. 1. FT-IR spectra of guanosine nucleotides and their platinum complexes: (a)  $\text{NH}_4\text{GpG}$ , (b)  $\text{cis-}[\text{Pt}(\text{NH}_3)_2(5'\text{-GMP})_2]^{2+}$ , (c)  $\text{cis-}[\text{Pt}(\text{NH}_3)_2(3'\text{-GMP})_2]^{2+}$ , (d)  $5'\text{-GMPNa}_2$ , (e)  $\text{cis-}[\text{Pt}(\text{NH}_3)_2(\text{GpG})]^+$ .

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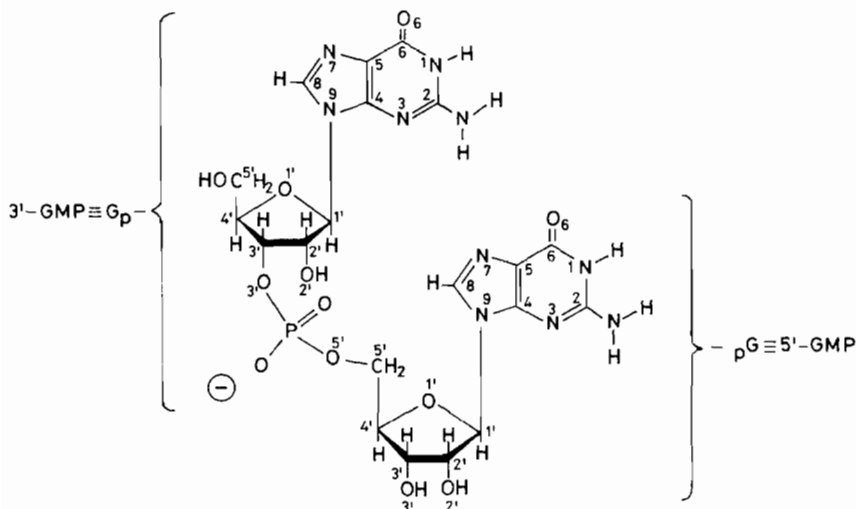


Fig. 2. Structural relationships in  $\text{NH}_4\text{GpG}$ , 3'-GMP, and 5'-GMP.

The FT-IR spectra of  $\text{NH}_4\text{GpG}$  shows marker bands at 802 and 797  $\text{cm}^{-1}$ . The spectra of  $\text{cis}[\text{Pt}(\text{NH}_3)_2(5'\text{-GMP})_2]^{2+}$  and  $\text{cis}[\text{Pt}(\text{NH}_3)_2(3'\text{-GMP})_2]^{2+}$  show these bands at 798 and 793  $\text{cm}^{-1}$ , these bands are assigned to the C3'-endo, anti, gg sugar-phosphate vibrations as has been reported previously [2]. However, the spectra of 5'-GMPNa<sub>2</sub>, having a C2'-endo, anti, gg conformation, shows the marker band at 822  $\text{cm}^{-1}$  but the characteristic band at 800  $\text{cm}^{-1}$  is absent. As is shown in Fig. 2 the structure of  $\text{NH}_4\text{GpG}$  consists of 3'-GMP(Gp-) and 5'-GMP(-pG) fragments. The two bands of  $\text{NH}_4\text{GpG}$  at 802 and 797  $\text{cm}^{-1}$  are thus assigned to ribose (-pG) and ribose (Gp-), respectively. Therefore, both sugar rings adopt predominantly a C3'-endo conformation. In aqueous solution, an NMR conformational analysis can be obtained by calculating the percentage of C2'-endo conformer of the dinucleotides by taking the value of  $10J_{1'2'}$  [6]. From the coupling constants (5.87 ppm,  $J_{1'2'} = 4.8$  Hz and 5.78 ppm,  $J_{1'2'} = 4.4$  Hz) in  $\text{NH}_4\text{GpG}$  the percentage of C2'-endo of both ribose rings were 48% and 44%, respectively, in aqueous solution. These results are different, however, from those of FT-IR spectroscopy and X-ray analysis of GpC salts in the solid state.

The FT-IR spectra of  $\text{cis}[\text{Pt}(\text{NH}_3)_2(\text{GpG})]^+$  with N7-N7 chelation in the GpG sequence shows a marker band at 800  $\text{cm}^{-1}$ , assigned to the C3'-endo and a new shoulder band at  $\sim 820$   $\text{cm}^{-1}$  related to a C2'-endo ring pucker. The former band will be related to the sugar-phosphate vibrational band of the ribose ring in (Gp-) and the latter is undoubtedly due to that of the ribose in (-pG). The band which was observed at 802  $\text{cm}^{-1}$  in  $\text{NH}_4\text{GpG}$  corresponding to the (-pG) ribose fragment has disappeared in  $\text{cis}[\text{Pt}(\text{NH}_3)_2(\text{GpG})]^+$  and a new shoulder band has been observed at 820  $\text{cm}^{-1}$  in the GpG-platinum

complex. It seems that the C2'-endo sugar-phosphate vibration in the (-pG) fragment is the same as that in 5'-GMPNa<sub>2</sub>.

In aqueous solution, the <sup>1</sup>H NMR spectra of the GpG-platinum complex exhibited one singlet at 6.04 ppm and one doublet at 5.87 ppm ( $J_{1'2'} = 6.9$  Hz) in agreement with results previously reported [3]. The absence of the  $J_{1'2'}$  value shows that the ribose ring adopts a 100% C3'-endo conformation, whereas the other  $J_{1'2'}$  value shows that the second ribose ring has a predominantly ( $\sim 70\%$ ) C2'-endo conformation. These <sup>1</sup>H NMR results are in agreement with the FT-IR spectroscopic data. It is found that the ribose conformation is both (Gp-) 797  $\text{cm}^{-1}$  and (-pG) 802  $\text{cm}^{-1}$  is C3'-endo, anti, gg in  $\text{NH}_4\text{GpG}$  and changes into (Gp-) 800  $\text{cm}^{-1}$  C3'-endo, anti, gg and (-pG) 820  $\text{cm}^{-1}$  C2'-endo, anti, gg, respectively, when a platinum atom is chelated to N7-N7 in the GpG sequence. This finding may be significant to the antitumor activity of cis-platinum and its interaction with DNA.

In conclusion, the conformational study of  $\text{NH}_4\text{GpG}$  and its adduct with cis-platinum reveals the following interesting sugar-phosphate change on complexation of the dinucleotide. In the natural free dinucleotide the sugar ring pucker in (Gp-) and (-pG) fragments is C3'-endo, anti, gg, however on complexation with the drug cis-platinum the (Gp-) sugar ring does not change, whereas the (-pG) sugar ring pucker changes predominantly into C2'-endo, anti, gg.

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