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**Conformations in crystals and solutions of d(CACGTG), d(CCGCGG) and d(GGCGCC) studied by vibrational spectroscopy**

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**ABSTRACT**

Crystals of self complementary DNA hexamers d(CACGTG), d(CCGCGG) and d(GGCGCC) were grown by vapour diffusion technique and studied by microRaman and microIR spectroscopies. The oligonucleotides were studied in parallel in solution by vibrational spectroscopy. A B→Z transition was detected by Raman spectroscopy during the crystallization process for d(CACGTG). Vibrational spectroscopy shows that the d(GGCGCC) crystals adopt a B geometry. On the contrary the d(CCGCGG) sequence which is shown to be able to undergo in solution or in films quite easily the B→Z transition, remains trapped in crystals in a geometry which may correspond to an intermediate conformation often proposed in models of the B→Z transition. The crystals used in this study were characterized by X-ray diffraction. The unit cell and space group have been determined.

**INTRODUCTION**

Fiber diffraction studies gave the first evidence of DNA polymorphism due to sequence and environmental conditions. In the last few years single crystal X-ray structures have shown that DNA helices are variable even in the same conformational family and many molecular details are now known, but difficulties persist to obtain information about some sequences due to the problem of crystallizing them (for review see 1-3). The results of X-ray crystal diffraction studies can be correlated with those obtained by other techniques and in particular by vibrational spectroscopy. A major advantage both of Raman and IR spectroscopies is the possibility to study oligo or polynucleotides under a wide variety of physical states: solutions, hydrated films or fibers, crystals. Thus for example the identity of the structures of the d(C-G)<sub>3</sub> hexamer in the crystal and of high salt poly d(G-C) in solution was obtained by Raman spectroscopy (4). Several Raman studies of crystal conformations have been already presented (5-9). On the contrary microIR studies of crystallized oligonucleotides have not been published up to now. We report here results obtained by micro IR and micro Raman spectroscopies on three crystals d(CACGTG), d(GGCGCC) and d(CCGCGG). In parallel these oligonucleotides have been studied in solution by vibrational spectroscopy. The crystallized oligonucleotide d(CACGTG) has been shown very recently by X-ray diffraction to adopt a Z geometry (10). We present here the Raman study of this crystal, allowing us to confirm the marker lines of left-handed A-T

base pairs. The results are in excellent agreement with previously proposed marker lines in the case of poly d(A-C).poly d(G-T) or poly d(A-T), but obtained in particular ionic strength conditions, in presence of nickel ions. The comparative study by Raman spectroscopy of the d(CACGTG) crystal and solution shows that the hexamer is in B form in solution and in Z form in the crystal. The d(GGCGCC) sequence was considered as a good candidate for a potential B geometry in crystalline state. The reverse 5'->3' sequence d(CCGCGG), should theoretically be able to adopt a Z geometry. Base pairs out of purine-pyrimidine alternance were used so as to avoid the formation of concatemers. Infrared spectra of d(CCGCGG) solutions and hydrated films show that this oligonucleotide is able to undergo quite easily the B->Z transition either by increase of the ionic strength (solutions) or by decrease of the relative humidity (films). However microRaman and microIR results obtained on the d(CCGCGG) crystal may reflect an intermediate structure between B and Z geometries.

#### MATERIALS AND METHODS

##### Synthesis and crystallization

The oligonucleotides d(CACGTG), d(CCGCGG) and d(GGCGCC) were synthesized by the phosphotriester method in solution from dimers with trisopropylsulfonyl nitrotriazole as the coupling reagent (11). They were purified by Sephadex G-25 gel column chromatography and subsequently by preparative high-performance liquid chromatography on reverse-phase Zorbax OMS 9.3 mm column. Crystals were grown using the vapor diffusion method. The oligonucleotide d(CACGTG) was crystallized in standard conditions, both in presence and absence of spermine (10). On the other hand the sequences d(CCGCGG) and d(GGCGCC) were crystallized with counterions not previously used in oligonucleotide monocrystals. The counterions employed with d(CCGCGG) were alaninamide, isopropylamide and serinol, while d(GGCGCC) has been crystallized both in standard conditions and with the basic counterions Ac(Arg)<sub>2</sub>NHET.

##### Characterization of crystals by X-ray diffraction

Crystals of the sequence d(CACGTG) had two different cell dimensions depending on the presence or absence of spermine in the crystallization solution. Both types of crystals examined by still and precession X-ray diffraction photographs were found to have the same orthorhombic space group P2<sub>1</sub>2<sub>1</sub>2<sub>1</sub>. Cell dimensions were: a = 16.1, b = 29.7, c = 41.6 Å with spermine and a = 17.6, b = 31.1, c = 44.4 Å without spermine (10). All the d(CCGCGG) crystals obtained had a similar morphology, independently of the counterion present. They were tetragonal plates, always smaller than 0.2 x 0.2 x 0.05 mm. The precession photographs clearly indicated a tetragonal system with a P4<sub>22</sub> space group. However the c axis cell dimension showed two distinct values for different crystals grown under the same conditions. Complete data sets to 2.5 Å resolution were collected for the two kinds of crystals using a FAST area detector with synchrotron radiation at Daresbury. The unit cell parameters determined were a = b = 40.9, c = 35.6 Å and a = b =

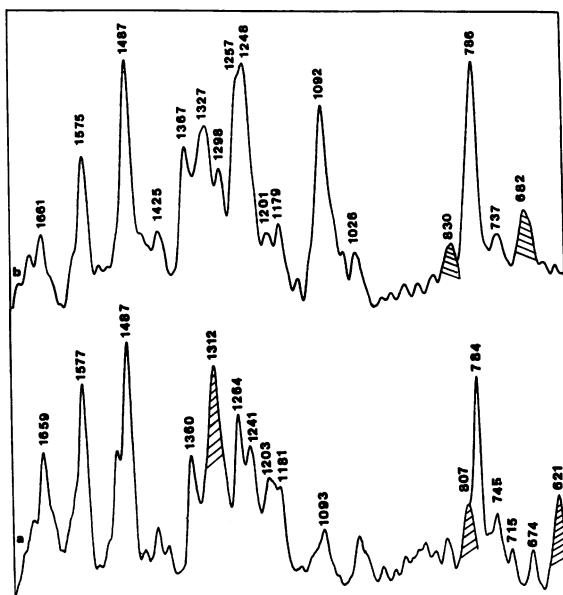


Figure 1 : Raman spectra of d(CACGTG) a : Crystal ; b : Solution ; marker lines characteristic of B form  and of Z form 

40.9,  $c = 31.2 \text{ \AA}$  respectively, therefore confirming the presence of two subtly different crystal forms. Crystals of the sequence d(GGCGCC) are thin plates. The precession photographs showed two 2-fold axis perpendicular to each other. The cell dimensions were  $a = 46$ ,  $b = 36$ ,  $c = 110 \text{ \AA}$ , one of the longest axis found for oligonucleotides. Precession photographs showed systematic absences for  $hk0$  when  $h + k = 2n + 1$  and also for  $00l$  when  $l = 2n + 1$ , thus the lattice must be centred (C) and the 2-fold axis parallel to the  $c$  axis must be a 2-fold screw axis, clearly indicating the space group  $C222_1$ . No sign of stacking was seen in the precession photographs. A complete data set to  $2.3 \text{ \AA}$  resolution was collected in a CAD4-Enraf Nonius with  $\text{Cu K}\alpha$  radiation. The accurate cell dimensions were  $a = 46.15 (1)$ ,  $b = 36.90 (2)$ ,  $c = 110.03 (3) \text{ \AA}$  and the space group  $C222_1$  was confirmed. From packing considerations three hexamers or eighteen base pairs should be placed in the asymmetric unit, approximately aligned along the  $c$  crystal axis though no clear relationship between neighbouring hexamers can be inferred at the present stage.

#### Raman spectroscopy

Samples were exposed to the 514.5 nm line from a Spectra-Physics model 2025 argon laser. The output power used was 400 mW at the source. Raman spectra were recorded with a Dilor Omars 89 multichannel spectrophotometer coupled to an IBM AT3 computer. The microscope used was an Olympus BH-2 model

with a 100 X objective. About 20mW was focused down to an area of 2-4 microns in diameter. Integration time was usually 10 s. Each spectrum is an average of about 250 integrations. Solvent background correction was performed by subtracting the solvent spectrum recorded in the same conditions.

#### FTIR spectroscopy

Infrared spectra were recorded using a Perkin Elmer 1750 spectrophotometer coupled to a PE 7700 minicomputer. Solution spectra were obtained by depositing a droplet of the oligomer solution between two ZnSe windows. DNA concentrations were around 2 OD/ $\mu$ l. Solution spectra were run at two different ionic strengths, 1M and 0.4M NaCl. Film spectra and relative humidity monitoring (H<sub>2</sub>O or D<sub>2</sub>O) were obtained as previously explained (12). Crystal transmission spectra were recorded using a Spectra-Tech. microscope attachment with mirror focussing. In this latter case an MCT detector was used. Data treatment included subtraction of the solvent spectrum, base line correction, smoothing following the Savitsky and Golay procedure (usually on 13 points).

### RESULTS AND DISCUSSION

#### d(CACGTG)

In figure 1 are shown the Raman spectra of d(CACGTG) in solution (1b) and in crystal (1a). The spectra are quite obviously very different reflecting two geometries for the oligonucleotide depending on its physical state. In the crystal the structure has been shown by X-ray diffraction studies to belong to the Z family (10). The Raman spectrum (1a) contains the marker lines both of AT and GC base pairs in a left-handed geometry, which we have previously characterized in the case of polynucleotides (13,14). We present in figure 2 with an expanded scale the Raman spectra of the left-handed geometries of poly d(G-C) (2a) and poly d(A-T) (2d) in two regions containing marker lines, i.e. in the base fingerprint region between 1400 and 1150 cm<sup>-1</sup> (left) and in the low wavenumber region between 850 and 600 cm<sup>-1</sup> (right). The spectrum of the d(CACGTG) crystal (2c) can be satisfactorily compared either with the experimental spectrum of Z form poly d(A-C).poly d(G-T) (13) or with a simulated spectrum obtained by a linear combination of Z form spectra of poly d(G-C) (2a) and poly d(A-T) (2d) taking into account the base composition of the oligonucleotide i.e. 2 GC base pairs per 1 AT base pair. The computed spectrum is presented in the same spectral regions in Figure 2b. The experimental spectrum of the d(CACGTG) crystal has the features of the Z geometries of both AT and GC base pairs. The line observed at 1360 cm<sup>-1</sup> can be assigned to contributions of guanosines and adenosines in a *syn* geometry, which have Raman lines respectively at 1354 and 1362 cm<sup>-1</sup> in the Z polymers poly d(G-C) and poly d(A-T). The strong line observed at 1312 cm<sup>-1</sup> is due to the overlap of the contributions of guanosines (strong line at 1314 cm<sup>-1</sup> in poly d(G-C)) and adenosines (lines located at 1332 and 1314 cm<sup>-1</sup> in poly d(A-T)). Similarly the strong line observed at 1264 cm<sup>-1</sup> in the crystal spectrum is due to a cytosine vibration detected at the same position in the case of Z poly d(G-C)(6).

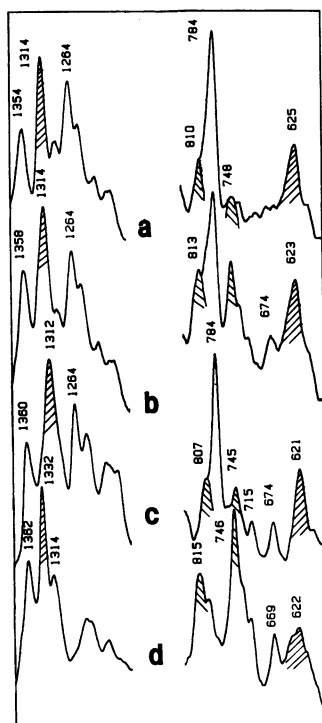


Figure 2 : Raman spectra in two conformation sensitive spectral regions ; left :  $1400-1200\text{ cm}^{-1}$  ; right :  $850-600\text{ cm}^{-1}$  of a) poly d(G-C) Z form (solution) ; b) computer spectrum 2a + 1d ; c) d(CACGTG) crystal ; d) poly d(A-T) Z form (solution) ; bases  $\text{////}$  backbone vibrations  $\text{\\\\\\\\}$

In the low wavenumber spectral region (between  $850$  and  $600\text{ cm}^{-1}$ ) the agreement between the experimental d(CACGTG) crystal spectrum (2c right) and the computed one (2b right) is also excellent. The experimental spectrum contains the main Z form marker line located at  $621\text{ cm}^{-1}$  involving a purine (guanine as well as adenine) breathing motion coupled through the glycosidic bond to a deoxyribose vibration. This mode is observed at  $621\text{ cm}^{-1}$  when the purine nucleotides are in a C3'-endo/*syn* geometry instead of  $682\text{ cm}^{-1}$  for guanosines and  $669\text{ cm}^{-1}$  for adenosines in C2'-endo/*anti* geometry (B form). The line observed at  $674\text{ cm}^{-1}$  in the crystal may be assigned to a thymidine vibration ( $669\text{ cm}^{-1}$  in poly d(A-T) Z form). The Z backbone modes in the crystal are observed at  $745$  and  $807\text{ cm}^{-1}$  at positions comparable to those of the same modes in the Z form polymers ( $748$  and  $810\text{ cm}^{-1}$  for poly d(G-C),  $746$  and  $815\text{ cm}^{-1}$  for poly d(A-T)). We must notice that this crystal is a relatively rare example of an oligonucleotide containing A-T base pairs without any chemical modifications such as methylation or bromination, which is found in a Z geometry by X ray diffraction (10). The Raman markers of Z

Table 1 : Main Raman lines sensitive to the B  $\rightarrow$  Z transition

B FORM				Z FORM					
Poly d(G-C)	Poly d(A-C) Poly d(G-T)	Poly d(A-T)	dCACGTG solution	Assignments	Poly d(G-C)	Poly d(A-C) Poly d(G-T)	Poly d(A-T)	dCACGTG crystal	Assignments
1363	1374	1374	1367	dA anti dG anti	1354	1354 1328 1314	1362	1360	dA, dG syn dG syn dA syn dA,dG syn
830	832	832	830	Backbone	810	804	815	807	Z Backbone
784	787	790	786		748	742	746	745	
682	730 682	727	737 682	dA anti dG anti	625	622	622	621	dG,dA syn

Poly d(G-C) from ref. (21), poly d(A-C).poly d(G-T) ref. (13), Poly d(A-T) ref. (14)

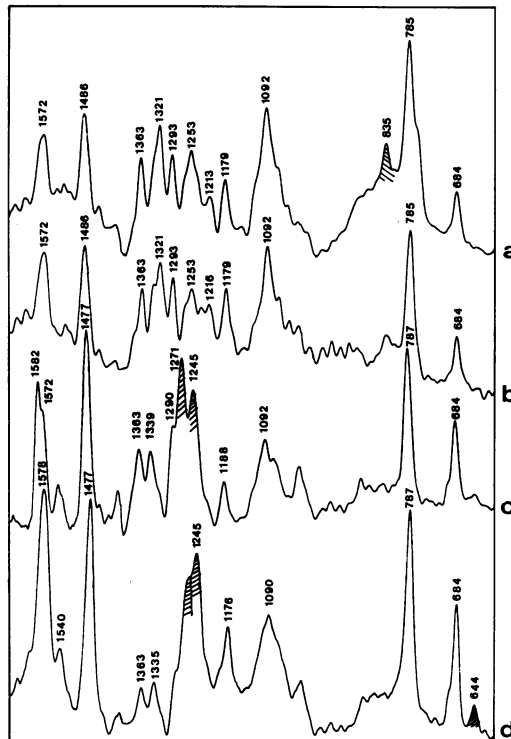


Figure 3 : MicroRaman spectra of crystals a) d(GGCGCC) + Ac(Arg)<sub>2</sub> NHET B form  $\text{////}$  ; b) d(GGCGCC), B form  $\text{////}$  ; c) d(CCGCGG) + alaminamide intermediate form  $\text{////}$  ; d) d(CCGCGG) + serinol intermediate form  $\text{////}$

form for A-T base pairs which we have detected here on the d(CACGTG) crystal spectrum confirm our previous assignments concerning the left-handed markers of A-T base pairs obtained in the case of polymers (poly d(A-C), poly d(G-T) and poly d(A-T)) but in particular experimental conditions i.e. in the presence of divalent metal ions (nickel) (13,14). The addition of these ions, which have been shown to stabilize the *syn* geometry of purines by interacting on the N7 site of the bases, and therefore the Z form of the sequences (15), does not modify the position of the Z form marker lines of A-T base pairs which we detect here for d(CACGTG) at the same positions without the presence of the metal ions.

The spectrum of d(CACGTG) obtained in solution (Fig. 1b) is quite different from the previously discussed one and clearly reflects another geometry. The Raman marker lines of a right-handed B form can be easily detected, as previously pointed out in the case of polynucleotides, oligonucleotides and native DNAs (16-19). In particular we observe a strong line at  $682\text{ cm}^{-1}$  reflecting the existence of guanines in C2'-endo/anti

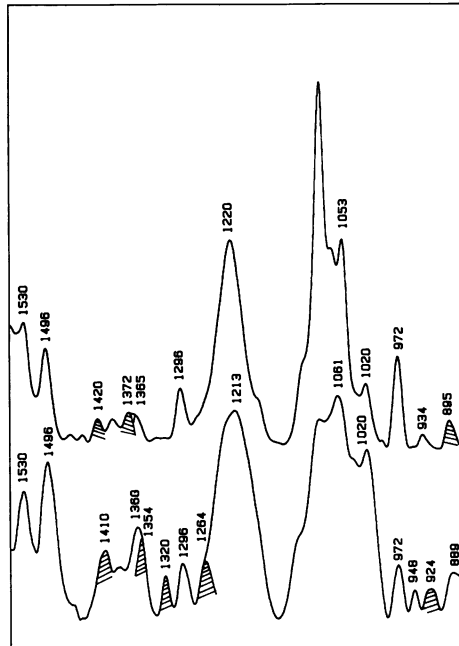




Figure 4 : FTIR spectra of d(CCGCGG) solutions : top : 0.4 M NaCl B form; bottom : 1 M NaCl Z form ; marker bands of B form  marker bands of Z form 

geometry, with a shoulder to the lower wavenumbers ( $669\text{ cm}^{-1}$ ) due to the thymidine contribution. The backbone vibrations are reflected by two lines located at  $830$  and  $786\text{ cm}^{-1}$ , the latter being superimposed on peaks due to cytosine as well as thymidine vibrations. In the base fingerprint region we detect the C2'-endo/anti guanosine vibration line at  $1367\text{ cm}^{-1}$ . All these results are summarized in table 1.

Thus the d(CACGTG) sequence crystallizes in a Z form while the B geometry is the conformation in solution. This type of effect had been first observed in the case of the d(C-G)<sub>6</sub> crystal which was obtained from a solution in which the hexamer was in B form (20). The structural change under the crystallization process has been relatively frequently observed, either reflected by a B→Z transition or by a B→A transition. However this is not systematically the case as will be shown by the study of the following oligonucleotide.

d(GGCGCC)

The sequence d(GGCGCC) has been studied by microRaman spectroscopy. It has been reported that sequences with initial purines and terminating pyrimidines will not go into the Z form (7). Thus this sequence was expected to be a good candidate for a still rare geometry of crystallized



oligonucleotides : B form. Figure 3 shows the Raman spectra of d(GGCGCC) crystallized in standard conditions (3b) and in presence of Ac(Arg)<sub>2</sub>NHET (3a). Both spectra are very similar and reflect a B form of the oligonucleotide : line at 835 cm<sup>-1</sup> due to the antisymmetrical phosphate stretching vibration in B geometry, strong marker line at 684 cm<sup>-1</sup> involving the guanosine breathing motion in a C2'-endo/anti geometry, characteristic profile in the 1200-1400 cm<sup>-1</sup> with positions and relative intensities of the lines identical to those previously published for B form d(C-G)<sub>n</sub> sequences (21).

#### d(CCGCGG)

Figure 4 shows the IR spectra of d(CCGCGG) solutions in 0.4 and 1M NaCl. They can be assigned using the marker IR bands of the different helices previously determined for poly d(G-C) (22) or d(CGCGGC) (23). The characteristic spectroscopic features of a B geometry are detected at low ionic strength and of a Z geometry at high ionic strength. The main spectral modifications observed under the B->Z transition of d(CCGCGG) are : (I) shift of a band located at 1420 cm<sup>-1</sup> assigned to a C2'-endo deoxyribose vibration to 1410 cm<sup>-1</sup> reflecting a C3'-endo sugar pucker; (II) shift of the band located at 1372 cm<sup>-1</sup> assigned to a ring vibration of the guanine coupled to a deoxyribose motion via the glycosidic linkage to 1354 cm<sup>-1</sup>, indicative of the anti->syn reorientation of the purine nucleotides; (III) presence of a new strong band at 1320 cm<sup>-1</sup> assigned to a syn geometry of the guanosine residue; (IV) presence of a band at 1264 cm<sup>-1</sup> (shoulder on the strong phosphate band) also due to a guanosine vibration; (V) shift of the phosphate antisymmetric stretching vibration to 1213 cm<sup>-1</sup> ; (VI) drastic decrease of the relative intensity of the absorption located at 1086 cm<sup>-1</sup> assigned to the symmetric stretching vibration of the phosphate groups; (VII) enhancement of the relative intensities of the 1061 and 1020 cm<sup>-1</sup> bands; (VIII) presence of a band at 924 cm<sup>-1</sup> characteristic of left-handed helices and (IX) decrease of the relative intensity of the band at 895cm<sup>-1</sup> characteristic of right-handed helices.

The B->Z transition of this oligonucleotide can also be induced in films by varying the water content of the sample. The figure 5 shows the IR absorption spectra of d(CCGCGG) films exposed to solutions of saturated salts in D<sub>2</sub>O corresponding to high (98%) and low (58%) relative humidities. The results obtained on films exposed to H<sub>2</sub>O saturated salts are identical to those obtained for solutions and the corresponding spectra will not be presented here. The decrease of RD induces in the 1750-1550 cm<sup>-1</sup> region shifts of the two bands located at 1681 and 1649 cm<sup>-1</sup> to lower wavenumbers (respectively 1666 and 1635 cm<sup>-1</sup>) similarly to what had been previously observed for poly d(G-C) (12), characteristic of a B->Z transition for the sequence. In the low wavenumber region between 1000 and 750 cm<sup>-1</sup> a decrease of the relative intensity of the 892 cm<sup>-1</sup> band and the emergence of a strong band at 927 cm<sup>-1</sup> reflect the B->Z transition induced by the decrease of the RD. All these modifications show that the d(CCGCGG) sequence is able

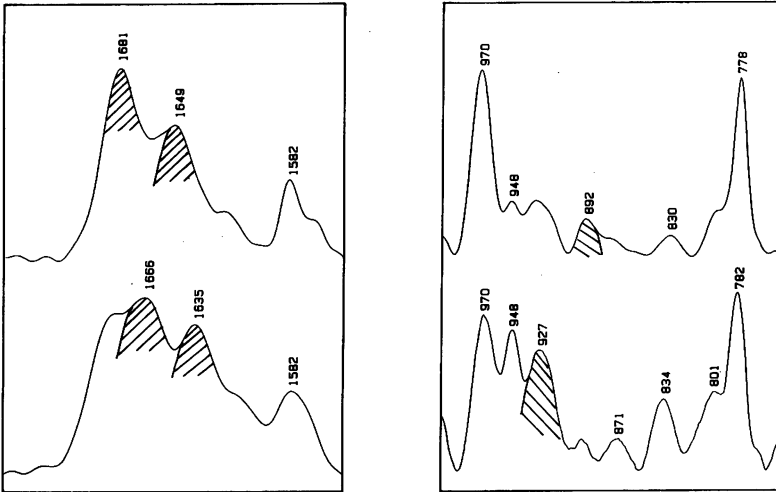


Figure 5 : FTIR spectra of d(CCGCGG) films exposed to D<sub>2</sub>O saturated solutions. Left between 1750 and 1550 cm<sup>-1</sup>. Right between 1000 and 750 cm<sup>-1</sup>. Top : high RH, B form. Bottom : low RH, Z form. Bases backbone vibrations

to undergo a right->left handed helical transition in solution as well as in films.

We have crystallized this oligonucleotide in presence of various counterions and the crystals have been studied by Raman and FTIR microspectroscopies. The crystals are not in a Z geometry. The figure 3 shows

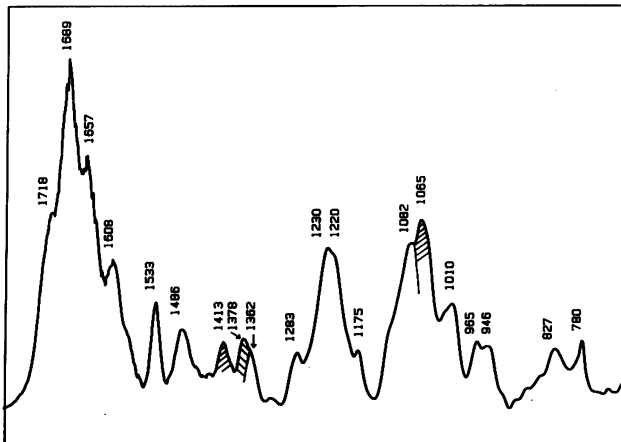


Figure 6 : Micro FTIR spectrum of d(CCGCGG) crystal + isopropylamine intermediate geometry between B and Z forms. Z genus B genus

the Raman spectra of the crystals obtained in presence of alaninamide (3c) and serinol (3d). The absence of a line around  $625\text{cm}^{-1}$  characteristic of guanosines in a C3'-endo/syn geometry and in contrast the existence of an intense line at  $684\text{ cm}^{-1}$  reflecting a C2'-endo/anti conformation of the purine nucleosides clearly shows that the structure does not belong to the left-handed Z form family. These spectra have mainly B form family characteristics ; however several points show that the structure which is studied does not belong to a classical B geometry. Thus we observe a decrease of the relative intensity of the phosphate line located at  $835\text{ cm}^{-1}$  in B geometry, an enhancement of a line at  $644\text{ cm}^{-1}$  and a very intense doublet at  $1245\text{-}1271\text{ cm}^{-1}$ . These three lines have been assigned to cytosine vibrations (7,16) and their intensities increase when destacking of the bases occurs (24). Such strong intensities of the  $1245\text{-}1271\text{ cm}^{-1}$  lines have been observed for a crystal of d(CpG) where base stacking should not be as important as in the case of longer oligonucleotides (7).

The infrared spectrum of a d(CCGCGG) crystal with isopropilamine (Figure 6) is different from the previously discussed B form spectrum recorded in solution or in films. Several absorptions involving the deoxyribose are affected. In particular we detect a strong band at  $1065\text{ cm}^{-1}$ , an important enhancement of the deoxyribose contribution at  $1010\text{ cm}^{-1}$ , a new band at  $946\text{ cm}^{-1}$  and a shift of the band located at  $1420\text{ cm}^{-1}$  to  $1413\text{ cm}^{-1}$  which would indicate a C3'-endo pucker. A decrease of the  $894\text{ cm}^{-1}$  band relative intensity is also detected. The comparison of the crystal spectrum (figure 6) and of the B and Z form solution spectra (figure 4) shows that the guanosines remain in an anti geometry (band at  $1378\text{ cm}^{-1}$ , no bands at  $1320$  and  $1264\text{ cm}^{-1}$ ).

It may thus be proposed that the microRaman and microIR spectra of d(CCGCGG) crystals obtained in presence of these different cations reflect a modified B geometry, may be an intermediate step in the B->Z transition. Such an intermediate step in the case of d(C-G)<sub>n</sub> sequences has been proposed after CD experiments. In ethanolic solutions, induction of the B->Z transition by increase of temperature allows to isolate an intermediate form characterized by its particular CD spectrum (25). Similarly a B\* form induced by increase of the ionic strength has been interpreted in terms of unfolding or premelting of the structure (26). Intermediates in the B->Z transition have also been detected by Raman spectroscopy. Spectra recorded at regular time intervals after addition of salt show the emergence of a Raman line around  $640\text{ cm}^{-1}$  prior to that of the characteristic Z form marker at  $625\text{ cm}^{-1}$  (21). The spectrum recorded in 1.5 M NaCl does not present the  $625\text{ cm}^{-1}$  line but clearly the existence of a line at  $640\text{ cm}^{-1}$  (27).

The d(CCGCGG) sequence which is in B form in solution at low ionic strength was expected to adopt a Z form in crystal, as the B->Z transition of this oligonucleotide is easily induced either by increase of the ionic strength in solution or by decrease of the hydration in films. Rather surprising is the result that d(CCGCGG) crystals are detected by micro IR and micro Raman neither in a classical B form nor in Z geometry. It seems thus

that the expected B->Z transition which should have occurred by crystallization has possibly been stopped by the presence of the counterions used in the crystallization process: isopropilamine, alaninamide or serinol.

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