# Normal coordinate analysis of 2'-deoxythymidine and 2'-deoxyadenosine

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Abstract. The proposed valence force field allows us to reproduce the vibration modes of 2'-deoxythymidine and 2'-deoxyadenosine. The present calculations are based on the Wilson GF-method and a non-redundant set of symmetrical coordinates. The calculated wavenumbers have been compared to the available Raman and infrared peak positions observed in solid, amorphous or aqueous samples. Moreover, the results obtained with the present force field allow us to assign some of the characteristic vibration modes for the thymidine and adenosine residues involved in DNA double-helical chains.

**Key words:** 2'-deoxythimidine, 2'-deoxyadenosine, DNA, vibration modes

## Introduction

We have recently published the normal coordinate analysis of 5'-dGMP (Ghomi and Taillandier 1985) and 2'-deoxycytidine (2'-dC) (Letellier et al. 1986). The non-redundant force field obtained in these calculations was also able to reproduce the vibrational characteristics of the cytidine (dC) and guanosine (dG) residues involved in DNA double helical chains (Letellier et al. 1986).

New spectroscopic studies on oligonucleotides and polynucleotides show that adenosine (dA) and thymidine (dT) residues may exist in both right-and left-handed conformations (Thomas and Peticolas 1983; Benevides et al. 1984; Thomas and Benevides 1985; Jollès et al. 1985; Adam et al. 1986 a, b). Moreover, some of the characteristic vibration modes of the dT and dA residues are found to be altered by both right → right and right → left conformational transitions. It would therefore be help-

ful to interpret these experimental phenomena by a normal coordinate treatment.

In this paper, our purpose is to obtain a reliable force field for 2'-deoxythymidine (2'-dT) and 2'-deoxyadenosine (2'-dA) for which recent Raman and infrared data are available. Obviously, the present calculations constitute the first step in the vibration mode analysis of the dT and dA residues encountered in oligonucleotides and polynucleotides.

#### Calculations

The normal coordinate analysis detailed here has been made using the Wilson GF-method (Wilson et al. 1955). The calculation formalism and the notations have been described in our former papers (Ghomi et al. 1985; Letellier et al. 1986).

To perform the numerical calculations, the original NCTB program (Shimanouchi 1968) has been vectorized in order to improve its run time on a Cray-1s computer. The array processing considerably accelerates the numerical calculations in comparison with the classical scalar computers.

As the out-of-plane modes of the pyrimidine and purine bases, particularly those of the thymine and adenine residues have been studied recently (Letellier et al. 1987a, b), our attention in this paper was especially focused on the planar modes of the bases. On the basis of these considerations, one expects 78 vibration modes from 2'-dT and 75 from 2'-dA. 99 internal coordinates are numbered in both 2'-dT (32 stretchings + 57 bendings + 10 torsions) and 2'-dA (33 stretchings + 57 bendings + 9 torsions). So, there are 22 redundant coordinates in 2'-dT and 24 in 2'-dA. To preserve the harmonic approximation of the potential field, these redundant coordinates have been removed by a standard  $B \cdot \bar{B}$  matrixproduct diagonalization procedure (Gusoni and Zerbi 1968) in order to obtain as many internal co-

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ordinates as the expected vibration modes (normal coordinates). As in the case of 5'-dGMP, the local symmetry of the different molecular sites has been taken into consideration and the redundancies have been resolved in the bases, sugars and C5'-tetrahedral sites separately (Ghomi and Taillandier 1985).

To construct *B* and *G* matrices the crystal structure of 2'-dT (Young et al. 1969) and that of 2'-dA (Watson et al. 1965) have been used. Both of the nucleosides are in *anti* position and contain a *C3'-exo* sugar pucker.

The elements of the interatomic interaction matrix, F, have been evaluated by a valence force field approximation. For the thymine-residue, we have used the uracil force constants (Ghomi et al. 1986) as initial values and introduced the methylgroup force field (Susi and Ard 1975) in the C5position and also the supplementary force constants which taken into account the  $H \rightarrow C$  replacement in the N1-site. Then, the force constants have been slightly refined around their initial values in order to obtain a good agreement for 2'-dT and its N3deuterated analog wavenumbers (Table 1a). As far as the adenine-residue is concerned, the stretching and bending force constants; i.e. diagonal terms of the F matrix, are those adopted by Majoube (1985) in his recent normal coordinate analysis. Unfortunately, non-diagonal valence force constants were not available and our preliminary calculations with only Majoube's diagonal force constants gave a very poor agreement with the experimental results especially with those arising from the C8-deuterated analog of 2'-dA.

We have therefore introduced a reasonable set of interaction force constants (Table 1b) and refined them in order to improve the agreement between the experimental and calculated results. Sugar force constants are those used in our previous calculations on 5'-dGMP and 2'-dC (Ghomi and Taillandier 1985; Letellier et al. 1986).

Cartesian displacements of the vibrating atoms, i.e. elements of the  $L_x$  matrix, have been transferred to a DPS 6/96 CII-Honeywell Bull computer connected to a SE-293 BBC plotter in order to draw the molecular deformations during the selected characteristic vibration modes.

#### Results and discussion

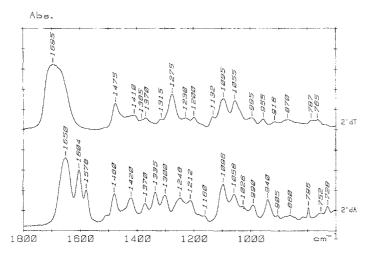
FT-IR and off-resonance Raman spectra of 2'-dT  $(\lambda_{\rm exc.} = 514.5 \text{ nm})$  in the solid state, have been performed by Mathlouthi et al. (1984) and resonant Raman spectra of 2'-dT and 2'-dA  $(\lambda_{\rm exc.} = 257 \text{ nm})$  in solution (pH 7) studied by Jollès et al. (1985). The

**Table 1 a.** Force constants of the thymine-residue. Bondstretch and stretch-stretch interaction force constants are in mdyne/A. Angular bending, bend-bend interaction and torsion force constants in mdyne/A. Stretch-bend interaction force constants are in mdyne. The notations and symbols are those used by Susi and Ard (1975). (a) Values used in our former calculations on uracil and its <sup>18</sup>O-derivatives (Ghomi et al. 1986); (b) Present work; (c) Susi and Ard (1975)

Bond-stretc		agonal for	ce constants Angular ben	ding F.C.	
BOND-STREET	1	/h l	, Angora Den	(a)	(b)
	(a) ;	(b)	C1'-N1	10/	1.909
C1'-N1	; ;	3.96	LT -NT		. 1.303
N-H	5.568	5.568	N−H	0.412	0.403
C-H	5, 292	5.292	С-н	0.409	0.475
C-N	6.38	6.28	C=0	1,213	1.213
C-C	5,202	6.37		1	; ;
C=0	11.00	10.70	<b>0</b> =¢)	1.647	1.957
C=C	9.00	9.00	R—C	0.521	0.621
	 		R—N	0.808	1.008
	. c.		'		:
C5CH3	0.10			,	;
		lagonal for	ce constants		
Str. bend	,		Str. Str.	Interactions	,
	(a)	(b)		(a)	(6)
	0.514	0.90	ortho	1.044	1.200
$\bigcirc$	; ;		, meta	-0.356	-0.356
~	; ;		para	0.557	0.557
	0.421	0.421	C=0,C=C	-0.225	-0.226
$\checkmark$	: :		C=0,ring	1.391	1.191
	0.177	0.197			
~	<u>;                                    </u>				
	Me t	hyl-group f	orce constant:	,	
	(c)	(b)		(c)	(6)
C5-CH3 st.	4,889 ± 0.178	5.189	C5-CH3	1.014 ± 0.078	1.014
Symm. st.	4.956 ± 0.034	4.956	Symm. bend.	0.569 ± 0:009	0.445
Asymm. str.	4.746 ± 0.032	4.746	Asymm. bend.	0.538 ± 0.009	0.405
	, ,	:	Rocking	0.663 ± 0.021	0.493
C-CH3 Str., Ring str.	0.471 ± 0.115	0.471	C-CH3 Str., Symm. bend	-0.465	-0.465

Table 1 b. Non-diagonal force constants of the adenine-residue. The diagonal force constants are those used by Majoube (1985) in his normal coordinate analysis of the adenine-residue (py): pyrimidic ring; (im): imidazolic ring

Non-diagonal force constants SirSir. F.C. (mdyn/A) Strbend. F.C. (mdyn) C=C,C-C(py) 0.500 C-N,DCH(py) 0.200 C-C,C=N(py) 0.500 C=N,DCH(py) 0.200 C=N,C-N(py) 0.551 C=C,DR (py) 0.810 C=C,C-N(py) 0.500 C-C,DR (py) 0.448 CC,CN(im,py) 0.600 C-N,DR (py) 0.531 C5-N6,ring 0.501 C5-N6,DR (py) 0.631 C8=N7,C2=N3 -0.100 C=N,DCH(im) 0.080 C8=N7,C4=C5 0.100 C-N,DCH(im) 0.400 C2=N3,C4=C5 0.200 C=C,DR(im) 0.530 Bendbend. F.C. (mdyn.A) C=N,DC(im) 0.531 N7CH,N3CH 0.300 C-N,DR(im) 0.530 N9CH,N3CH 0.290 CC,DR(im,py) 0.530 N9CH,N3CH 0.290 CC,DR(im,py) 0.260 N9CH,N3CH 0.290 CC,DR(im,py) 0.260 N9CH,N3CH -0.1501 CN,DR(im,py) 0.260	Adenine-residue									
C=C,C-C(py) 0.500 C-N,DCH(py) 0.200 C-C,C=N(py) 0.500 C=N,DCH(py) 0.200 C=N,C-N(py) 0.500 C=C,DR(py) 0.810 C=C,C-N(py) 0.500 C-C,DR(py) 0.448 CC,CN(lm,py) 0.600 C=N,DR(py) 0.322 CN,CN(lm,py) 0.600 C-N,DR(py) 0.531 C6-N6,rlng 0.501 C6-N6,DR(py) 0.431 C8=N7,C2=N3 -0.100 C=N,DCH(lm) 0.080 C8=N7,C4=C5 0.100 C-N,DCH(lm) 0.400 C2=N3,C4=C5 0.200 C=C,DR(lm) 0.530 Bend,-bend, F.C. (mdyn.A) C=N,DR(lm) 0.631 N7CH,N3CH 0.300 C-N,DR(lm) 0.530 N7CH,N3CH 0.300 C-N,DR(lm) 0.530 N7CH,N3CH 0.290 CC,DR(lm,py) 0.530	,									
C-C, C=N(py) 0.500 C=N, DCH(py) 0.200 C=N, C-N(py) 0.551 C=C, DR (py) 0.810 C=C, C-N(py) 0.550 C-C, DR (py) 0.448 CC, CN(lm, py) 0.600 C=N, DR (py) 0.531 C6-N6, ring 0.501 C6-N6, DR (py) 0.531 C6=N7, C2=N3 -0.100 C=N, DCH(lm) 0.080 C8=N7, C4=C5 0.100 C-N, DCH(lm) 0.400 C2=N3, C4=C5 0.200 C=C, DR (lm) 0.530 Bend, -bend, F.C. (mdyn, A) C=N, DR (lm) 0.531 N7CH, N3CH 0.300 C-N, DR (lm) 0.530 N3CH, N3CH 0.290 CC, DR (lm) 0.530 N3CH, N3CH 0.290 CC, DR (lm) 0.530	StrStr. F	.C. (mdyn/A)	Strbend. F.	C. (mdyn)						
C=N, C-N(py) 0.651	C=C,C-C(py)	0.500	с-и, осн (ру)	0.200						
C=C,C-N(py) 0.500 C=C,DR(py) 0.448  CC,CN(lm,py) 0.600 C=N,DR(py) 0.322  CN,CN(lm,py) 0.600 C=N,DR(py) 0.631  C6-N6,rlng 0.501 C6-N6,DR(py) 0.431  C8=N7,C2=N3 -0.100 C=N,DCH(lm) 0.080  C8=N7,C4=C5 0.100 C=N,DCH(lm) 0.400  C2=N3,C4=C5 0.200 C=C,DR(lm) 0.530  Bendbend. F.C. (mdyn.A) C=N,DR(lm) 0.531  N7CH,N3CH 0.300 C-N,DR(lm) 0.531  N7CH,N3CH -0.1210 C1'N9,DR(lm) 0.530  N9CH,N3CH 0.290 CC,DR(lm,py) 0.260	C-C, C=N(py)	0.500	С=N, ДСН(ру)	0.200						
CC, CN(lm, py) 0.600 C=N, DR (py) 0.322 CN, CN(lm, py) 0.600 C=N, DR (py) 0.631 C6=N6, r lng 0.501 C6=N6, DR(py) 0.431 C8=N7, C2=N3 -0.100 C=N, DCH(lm) 0.080 C8=N7, C4=C5 0.100 C=N, DCH(lm) 0.400 C2=N3, C4=C5 0.200 C=C, DR(lm) 0.530 Bendbend. F.C. (mdyn.A) C=N, DR(lm) 0.631 N7CH, N3CH 0.300 C=N, DR(lm) 0.531 N7CH, N1CH -0.1210 C1'N9, DR(lm) 0.530 N9CH, N3CH 0.290 CC, DR(lm, py) 0.260	C=N, C-N(py)	0.651	C=C,DR (py)	0.810						
CN, CN (lim, py) 0.600	C=C,C-N(py)	0.500	C-C,DR (py)	0.448						
C6-N6, ring 0.501 C6-N6, DR(py) 0.431  C8=N7, C2=N3 -0.100 C=N, DCH(lm) 0.080  C8=N7, C4=C5 0.100 C-N, DCH(lm) 0.400  C2=N3, C4=C5 0.200 C=C, DR(lm) 0.530  Bend, -bend, F.C. (mdyn.A) C=N, DR(lm) 0.631  N7CH, N3CH 0.300 C-N, DR(lm) 0.531  N7CH, N1CH -0.1210 C1'N9, DR(lm) 0.530  N9CH, N3CH 0.290 CC, DR(lm, py) 0.260	CC, CN(Im, py)	0.600	C=N,DR (py)	0.322						
C8=N7, C2=N3     -0.100     C=N,DCH(lm)     0.080       C8=N7, C4=C5     0.100     C-N,DCH(lm)     0.400       C2=N3, C4=C5     0.200     C=C,DR(lm)     0.530       Bendbend. F.C. (mdyn.A)     C=N,DR(lm)     0.631       NZCH, N3CH     0.300     C-N,DR(lm)     0.531       NZCH, N1CH     -0.1210     C1'N9,DR(lm)     0.530       N9CH, N3CH     0.290     CC,DR(lm,py)     0.260	CN, CN(Im, py)	0.600	C-N, DR (py)	0.631						
C8=N7, C4=C5     0.100     C-N, DCH(lm)     0.400       C2=N3, C4=C5     0.200     C=C, DR(lm)     0.530       Bendbend. F.C. (mdyn.A)     C=N, DR(lm)     0.631       NZCH, N3CH     0.300     C-N, DR(lm)     0.531       NZCH, N1CH     -0.1210     C1'N9, DR(lm)     0.530       N9CH, N3CH     0.290     CC, DR(lm, py)     0.260	C5-N5,rlng	0.501	C6-N6, DR (py)	0.431						
C2=N3, C4=C5     0.200     C=C,DR(lm)     0.530       Bendbend. F.C. (mdyn.A)     C=N,DR(lm)     0.631       N7CH, N3CH     0.300     C=N,DR(lm)     0.531       N7CH, N1CH     -0.1210     C1'N9,DR(lm)     0.530       N9CH, N3CH     0.290     CC,DR(lm,py)     0.260	C8=N7, C2=N3	-0.100	C=N,DCH(lm)	0.080						
Bendbend. F.C. (mdyn.A)     C=N, DR(lm)     0.531       NZCH, N3CH     0.300     C-N, DR(lm)     0.531       NZCH, N1CH     -0.1210     C1'N9, DR(lm)     0.530       N9CH, N3CH     0.290     CC, DR(lm, py)     0.260	C8=N7,C4=C5	0.100	C-N,DCH(lm)	0.400						
N7CH, N3CH 0.300 C-N, DR(lm) 0.531 N7CH, N1CH -0.1210 C1'N9, DR(lm) 0.530 N9CH, N3CH 0.290 CC, DR(lm, py) 0.260	C2=N3,C4=C5	0.200	C=C,DR(lm)	0.530						
N7CH,N1CH -0.1210 C1'N9,DR(lm) 0.530 N9CH,N3CH 0.290 CC,DR(lm,py) 0.260	Bendbend.	F.C. (mdyn.A)	C=N,DR(lm)	0.531						
N9CH,N3CH 0.290 CC,DR((m,py) 0.260	илсн, изсн	0.300	C-N, DR((m)	0.531						
	NZCH, N1CH	-0.1210	C1'N9, DR(lm)	0.530						
N9CH,N1CH -0.1501 CN,DR(lm,py) 1.230	мэсн, мэсн	0.290	CC, DR(lm,py)	0.260						
	N9CH, N1CH	-0.1501	CN, DR(lm, py)	1.230						



**Fig. 1.** Infrared absorption spectra of the thin films of amorphous 2'-deoxythymidine (*up*) and 2'-deoxyadenosine (*down*)

infrared spectra of the thin films of amorphous 2'-dT and 2'-dA recorded in our laboratory are shown in Fig. 1 for the spectral region between 1800 and 700 cm<sup>-1</sup>. The above experimental data together with the 2'-dT infrared spectrum in  $D_2O$  (N3-deuterated) and that for the 2'-dA: C8-deuterated derivative (spectra not shown) have been utilized for adjusting the force constants as well as for comparison with the calculated results (Tables 2, 3 and 4).

## A. 2'-deoxythymidine and its N3-deuterated analog

Table 2 compares the calculated and experimental wavenumbers for 2'-deoxythymidine between 1800 and 650 cm<sup>-1</sup>. In Table 4, the comparison was made between the experimental and calculated infrared band-shifts (amorphous sample) upon N3-deuteration.

The majority of the infrared and Raman bands in the 1800-1270 cm<sup>-1</sup> region originate from the thymidine-residue. It is obvious that because of the crystalline field and Fermi-resonance effects the number of peaks observed in solid samples is larger than that predicted from the normal coordinate calculations on an isolated molecule. So, we have made our assignments on the basis of the experimental results arising from aqueous or amorphous samples. Between 1800 and 1600 cm<sup>-1</sup>, the vibrational spectra mainly arise from the C=O and C=C bondstretch modes. They give rise to an intense and nonresolved infrared band situated at 1695 cm<sup>-1</sup>, not affected by N3-deuteration. The intense infrared band situated around 1475 cm<sup>-1</sup> (Fig. 2) is that observed in poly d(A-T) · poly d(A-T) spectra for both right- and left-handed conformations (Adam et al. 1986 a, b). In contrast, the vibration mode around 1410 cm<sup>-1</sup> (Fig. 2) shifts from 1425 cm<sup>-1</sup> to 1401 cm<sup>-1</sup> in poly  $d(A-T) \cdot poly d(A-T)$  infrared spectra upon the  $B \rightarrow Z$  transition. The sugar-pucker CH2-scissoring mode situated around 1315 cm<sup>-1</sup> in 2'-dT (Fig. 2) is found to be altered by the right  $\rightarrow$  right transition of poly  $d(A-T) \cdot poly d(A-T)$  showing an infrared band-shift from 1327 cm<sup>-1</sup> (B form) to 1317 cm<sup>-1</sup> (A form). The intense infrared band around 1275 cm<sup>-1</sup> (Fig. 2) arises from the thymineresidue and is sensitive to N3-deuteration which makes it shift toward higher wavenumbers (Table 4).

The good agreement between the experimental and calculated wavenumbers situated below 1250 cm<sup>-1</sup>, corresponding mainly to the sugar vibration modes, confirms the validity of the 2'-deoxyribose force field. In Fig. 2 we have illustrated graphically the sugar modes responsible for the two intense infrared bands situated at 1095 and 1055 cm<sup>-1</sup> (Table 2).

The infrared band at 765 cm<sup>-1</sup>, also observed in polynucleotide spectra (Adam et al. 1986a, b), has been assigned to the thymine-residue out-of-plane vibrations by our very recent calculations (Letellier et al. 1987a).

# B. 2'-deoxyadenosine and its C8-deuterated analog

In Table 3, a comparison has been made between the experimental and calculated wavenumbers of 2'-dA in the 1800-700 cm<sup>-1</sup> spectral region. In Table 4 the calculated wavenumber-shifts are compared to the infrared band displacements upon C8-deuteration in the adenine residue.

On the basis of the calculated results, the infrared bands situated at 1650, 1604, 1578 and 1510 cm<sup>-1</sup> involve mainly the pyrimidic ring vibrations (Fig. 3). The last one gives rise to an intense Raman peak observed in the A and B forms of poly d (A-T) · poly d (A-T) (Thomas and Benevides

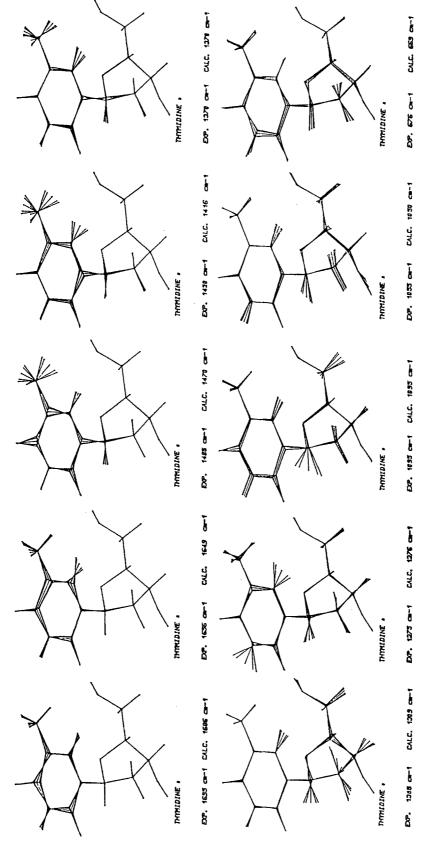


Fig. 2. Representation of the characteristic vibration modes of 2'-deoxythymidine. Experimental wavenumbers arising from the Raman or infrared spectra are also reported. See also Table 2

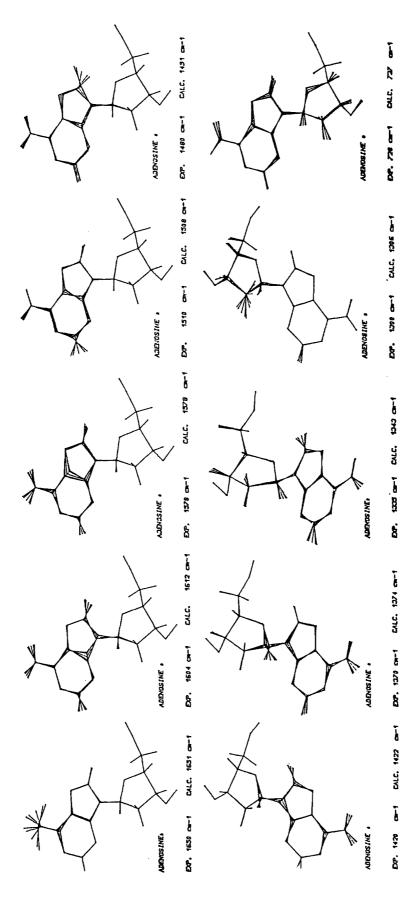


Fig. 3. Representation of the characteristic vibration modes of 2'-deoxyadenosine. Experimental wavenumbers arising from the infrared spectra of amorphous samples are also reported. See also Table 3

Table 2. Comparison between the experimental and calculated results for 2'-deoxythymidine. (a) The wavenumbers (cm<sup>-1</sup>) of 2'-dT neutral solution resonant Raman spectrum ( $\lambda_{\rm exc.} = 257$  nm) (Jollès et al. (1985); (b) Off-resonance Raman peak positions of the crystalline 2'-dT ( $\lambda_{\rm exc.} = 514.5$  nm) (Mathlouthi et al. 1984); (c) FT-IR peak positions of the crystalline 2'-dT ( $\lambda_{\rm exc.} = 514.5$  nm) (Mathlouthi et al. 1984); (d) I.R. wavenumbers of the thin films of amorphous 2'-dT (see also Fig. 1). The assignments are based on the internal coordinates for which the potential energy distribution (PED in per cent) is reported

to 200 to 100 to 100 to 100 to 100 to		CALCULATED		EXPERIME	EXPERIMENTAL		CALCULATED			
Ran	nan	IR		FREQ.	ASSIGNMENTS (PED %)	Raman	IR		FREQ.	ASSIGNMENTS (PED Z)
No. 200 201 100 100 10	ot play have THE bald you down you and		1705 d	1756	C2=02 (49);C4=04 (25)		۵.		1113	01'C4'H (35);01'C1'H (11); C5'C4'H (10);C1'-C2' (8)
		c	1695	:	C4=04 (35);C2=02 (19);C5=C6 (12)			1095		02-N3 (46);N3-04 (45);N4-06 (9) 02'04'H (44);O4'04'H (40)
1656	b 1666 b 1642	1665		1649	C5=C6 (41);C4=04 (21)	1068 1068	1070	d		04'C1'H (14);N1C1'H (13); C4'-C5' (9);C3'-03' (8)
a 1476	5 1486	1482	1475		C4-C5 (24) #N1-C2 (13)	1054	c	1055		C3'03'H (28);01'C1'H (12); C2'-C3' (11);C2-N3 (7)
	1460	1460		- - -		1028	1030			C1'C2'H (24);C3'-C4' (10); C3'-03' (7)
	1440 b	1440 c	d	:	CH3 - asymmetric deformation (85)	1020	c 1012		1022	C2'-C3' (16);C3'O3'H (12); O1'C1'H (8)
	1408 b	1415 E 1405		:	N1-C6 (24);C17-N1 (10);C5C6H (9); N1C6H (7)	b 1006	1012 c 1005	d 995		C3°C4'H (15);C5°C4'H (12); O5°C5'H (11);C5'-05' (8)
a	1392 b	1392	1385	1388	CH3 - symmetric deformation (39); CH7-NH (8)	b 976	975			CH3 - rockins
1372	1368	1365		:	CH3 - symmetric deformation (19); C2-N3 (18);C5-CH3 (8);N1-C6 (6)	, b 964	с 960	d 955		C5'-05' (14);C5'C4'H (11)
	ь 1326	1340 £		2	HC2'H (19);C4'-01' (10); C4'-C5' (9);C1'-01' (9)				:	05'C5'H (21);C4'C5'H (20); C3'C4'H (12);C5'-05' (19) C2'-C3' (9)
		1320		1309	HC2'H (38);C4'-01' (6);C4'-C5' (6)				: : 936	C5'-05' (28);C3'C2'H (16);
		1292		2	C4*-C5* (10);N3-C4 (9);N1C1*H (7); C5C4H (4)		c	ď		C1'C2'H (13)
	1282 1	1275		1276 1	CN3H (35) #C5-CH3 (14) #N1C6H (8) # C5C6H (6)	ь 902	910 c 900	916	2	C4'C5'H (29);05'C5'H (26); C4'-01' (9)
		1255		: 1255 : 1255	C3'-03' (13);C4'-01' (8); C1'-01' (7);HC3'03' (7)		8 <b>8</b> 3		, : 886 :	C37-037 (26);C37C27H (44); C57-057 (42);C27-C37 (9)
				: 1243	C5'05'H (64);HC5'H (7);C5'-05' (7)	b 874	672	d 870		C5-CH3 (16);C4-C5 (12)
3.	h	c	d		HC57H (52):057C57H (42)	ь	c		# 864 #	64'C3'H (37)%C2'C3'H (33)
1234	1232	1225		: 1222 :	CN3H (29);CS-CH3 (29);C2-N3 (4)	854	852		; 855 ;	C4'C5'H (29);05'C5'H (20); C4'-04' (18)
	1202	1200	1200 1200		C37037H (30)#C27-C37 (12)# C37-037 (9)#C37-C47 (8)				5	C2=02 (14);NI-C6 (12); C4-C5 (8);N1-C2 (8)
				: 1188 :	C17-C27 (11);04/C47H (10); C17-C17 (9);C27C17H (7)	5 794 5	792	787		01*-N1 (14);05-0H3 (7); N101*01* (7)
	ь 1176	c 1175		: 1184 :	03'C3'H (20);C4'-C5' (20); C3'-C4' (44);C2'C3'H (6)	738	734		: 725 :	C1'-01' (15);N38404 (8);
		_		5	C4'C5'H (16);C4'-C5' (13); O5'C5'H (10)	d			707	C3*C4*O1* (16) #C3*C2*H (14) # C2*C3*C4* (14)
	ь 1124	1125	1132	: 1133 :	N1-C2 (12);C4'C5'H (10); N3-C4 (10);C4-C5 (8)	676	672		: 669 :	62N964 (10);N36465 (9); 62'61'01' (9);61'01'64' (7)
				E					:	

1985). The 1480 cm<sup>-1</sup> infrared band has an imidazolic character and is sensitive to C8-deuteration. In polynucleotide infrared spectra this mode is superimposed on that arising from the thymine residue (Table 2, Fig. 2). The vibration mode at 1,420 cm<sup>-1</sup> is also shifted by C8-deuteration and is found to be altered by both the right → right and right → left-handed conformational transitions (Adam et al. 1986 a, b). As in the case of poly d (G-C) · poly d(G-C) (Ghomi et al. 1984) the vibration mode involving mainly the C1'-N9 bond-stretch vibration is taken to be one of the most important infrared markers for detecting the right → lefthanded transition of poly d(A-T) poly d(A-T). The present calculations assign this vibration mode to the 2'-dA infrared band situated at 1,370 cm<sup>-1</sup> (Fig. 3, Table 3).

Finally, the vibration mode at 1335 cm<sup>-1</sup> (Raman and infrared spectra) altered by C8-deuteration (Table 4) is found around 1340 cm<sup>-1</sup> in both right-and left-handed forms of poly d (A-T) poly d (A-T).

Below 1250 cm<sup>-1</sup>, the calculated results take account of the infrared modes arising from the sugar vibration modes.

It should be mentioned that the narrow infrared band situated at 796 cm<sup>-1</sup> is assigned to an adenine-residue out-of-plane vibration mode by our recent calculations (Letellier et al. 1987 b).

# Conclusion

As shown above, our non-redundant valence force field allows us to assign the most characteristic

Table 3. Comparison between the experimental and calculated results for 2'-deoxyadenosine. (a) Resonant Raman peak positions ( $\lambda_{\text{exc.}} = 257 \text{ nm}$ ) from 2'-dA neutral solutions (Jollès et al. 1984); (b) I.R. wavenumbers of the thin films of amorphous 2'-dA (see also Fig. 1). The assignments are based on the internal coordinates for which the potential energy distribution (PED in per cent) is reported

		RIMENTAL : CALCULATED		EXPERIMENTAL		: CALCULATED		
			ASSIGNMENTS (PED %)	Raman (a)	IR (b)	: FREG.	ASSIGNHENTS (PED %)	
163D (sh)	1604	: 1651 : 1642	HN6H (39);C6-N6 (24);C6N6H (22) C4-N9 (18);N3-C4 (18);C2≕N3 (9)		1058	: 1861 : 1861	NC2H (13)FC5-H7 (8)FC4-N9 (7)F C4=C5 (6)	
1584		157B	C5-C6 (31);C4=C5 (12);C5-N7 (6)			: 1053	C5'-05' (34);C5'05'H (15)	
		1508	NC2H (26);C2=N3 (15);N1-C2 (14); C5-C6 (12)		1026	: 1026	057C57H (13)#017C47H (10)#	
1484		1491	NC8H (27);N7=C8 (25);C2=N3 (12); N3-C4 (11)			: 1002	01'C1'H (10);C4'-01' (10) C1'-G2' (9);N9C1'H (7);	
		1452	N1=C6 (46):C6-N6 (11)			3	C3*C4*H (6)	
1424	1420	1422	C8-N9 (24);C1*-N9 (15)		990	: 999	C6N6H (46);N1=C6 (15);N3-C4 (7)	
	1370	1374	017-N9 (26);06-N6 (11)			976 8	C37-C47 (27);C37037H (23)	
	- 000	1343	NC8H (13);NC2H (7);C2=N3 (6)		;	: 954 :	C3'C4'H (22);C5'C4'H (14); C5'-05' (11)	
1332	1335	: 1324	C4"-C5" (16)#C4"-01" (14)			: 944	C3'C2'H (25);C5'+05' (9);	
	1300	1306	HC27H (48);C17C27H (11)		940	:	C3*C2*H (8)	
	1260 (sh)	1264	C1'-01' (16);C3'-03' (12)			: 931 :	C5-N7 (29) #N7=C8 (7)	
1248	1248	1244	HC57H (19);NC2H (10);N9C17H (9)			: 925 :	C4*C5*H (28);05*C5*H (17); C5*-05* (15)	
		1230	HC5'H (44);NC9H (9)		905	905	C4=C5 (16);C4-N9 (15)	
	1212	1215	NC8H (23)#N7=C8 (17)			# 871	C2'C3'H (37);C4'C3'H (30)	
		1206	037-047 (19);037037H (12); 047-057 (9)		860 .	: 861 :	C4'C5'H (25);C4'-01'(20); 05'C5'H (17)	
		1192	04'-05' (17);03'03'H (13); 01'-02' (11);02'01'H (10)			: 837 :	C3'03'H (32);C3'-03' (15)	
	1480 (sh)	: 1183	N1-C2 (31);C2=N3 (6)		815	# 813	C5'05'H (60);C5'-05' (9)	
	1160 (ah)		047057H (21);027-037 (15);		752	: 775	C17-N9 (14);C8-N9 (12)	
		: : 1436 :	C47-C57 (7) C31-C32 (20)3047C47h (12)% C47-C57 (12)		728	: 727	C4'-04' (12);C1'01'C4' (8); C5-N7 (7);C6-N6 (4);C4=C5 (3) C4-N3 (2);C8=N7 (2);N3=C2 (2) C4-N9 (2)	
1100 (sh)	1096	: : 1106 :	01'C1'H (48):C2'C1'H (47); 01'C4'H (41);C3'-03' (9)			714	C3'C4'01' (13);C2'C3'C4' (11); C3'C4'H (7)	
		1097	911047H (21):05105TH (19): 641-657 (7):011017H (6)			M M	C2N1C6 (15);N3-C4 (13);N3C4C5 (7)	
		1078	047-027 (40);047047H (7); 027-037 (7)					
		: 1067	N1+C2 (17);C6-N6 (45); C2N3C4 (44);N3-C4 (14)					

**Table 4.** Effect of deuteration in N3-site of the thymine residue in 2'-dT and in C8-site of the adenine-residue in 2'-dA. The calculated shift (in cm<sup>-1</sup>) is reported in parenthesis in front of each characteristic wavenumber. In the same manner the I.R. (amorphous samples) band shifts are indicated. — non-measurable, \*\* I.R. bands appearing after deuteration

2'	-deoxyth	ymidine		2'-deoxyadenosine				
exp.		cal.		exp.	exp.		cal.	
1695	/ )	1756	(0)	1650	(0)	1651	(0)	
1633	()	1686	(-2)	1504	(-1)	1512	(-8)	
1475	(0)	1470	(-1)	1578	(-3)	1578	(-5)	
	1440	1443	(0)	1510	()	1508	(-1)	
1410	(-5)	1415	(-1)	1480	(-17) **	1491	(-30)	
1385	(0)	1388	(-2)	;	1420	1452	(-29)	
1370	(0)	1370	(-2)	1420	(-15)	1422	(-7)	
	()	1325	(-1)	1370	(~3)	1374	(-1)	
1315		1309	(0)	1335	(-10)	1343	(-13)	
1275	(+25)	1275	(+24)	1300	(-2)	1305	(-1)	
1230	(+10)	. 1222	(+15)	1248	(-15)	1244	(-18)	
1200	(0)	1210	(+1)	1212	()	1215	(-232	
1132	(+18)	1133	(+9)	;				
1095	(+3)	1095	(+5)	10.96	(0)	10.97	(0)	
1055	()	1050	(-Z)	940	(-4)	944	(0)	
870	()	885	(-55)	728	(-5)	727	(-8)	

vibration modes of 2'-dT and 2'-dA. It is known that the conformation of both these nucleosides are modified when they are found in the right- and left-handed DNA double-helical chains. Consequently, the vibration modes associated with them will be altered by the conformational changes involved with the right  $\rightarrow$  right or right  $\rightarrow$  left-handed transitions. In order to interpret this kind of spectral evolution the presently proposed force field seems to be quite promising.

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