106.3 (3)

106.6 (4) 103.0 (3) 127.2 (4)

117.3 (4)

115.5 (3)

C6-N1-C2	114.4 (3)	C4—C3—C2
C6-N1-C5	114.8 (3)	C5-C4-C3
C2-N1-C5	104.8 (3)	C4-C5-N1
N1C2C3	104.2 (3)	01—C7—O2
N1-C2-C7	111.6 (3)	01—C7—C2

114.1 (3)

H atoms attached to N and O were located on a difference electron-density map and their coordinates refined with  $U_{iso} = 1.3U_{eq}$  of parent atom.

O2-C7-C2

Data collection: XSCANS (Siemens, 1993). Cell refinement: XSCANS. Data reduction: XSCANS. Program(s) used to solve structure: SIR92 (Altomare *et al.*, 1994). Program(s) used to refine structure: SHELXL93 (Sheldrick, 1993). Molecular graphics: SHELXTL/PC (Sheldrick, 1990). Software used to prepare material for publication: SHELXL93.

Financial support from DGAPA (UNAM) and CONACyT is gratefully acknowledged. WFR is indebted to NSERC, Canada. The authors want to thank R. Patiño, L. Velasco, I. Chávez and F. del Río for spectroscopic technical assistance.

Supplementary data for this paper are available from the IUCr electronic archives (Reference: BK1308). Services for accessing these data are described at the back of the journal.

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Acta Cryst. (1997). C53, 1692-1694

# *n*-Butyl $\alpha$ -Cyano-4-[2-(4-pyridyl)ethenyl]cinnamate†

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(Received 6 March 1997; accepted 26 June 1997)

#### Abstract

The crystal structure of the title compound,  $C_{21}H_{20}$ - $N_2O_2$ , is compared with related compounds. There are two crystallographically independent molecules, which form two respective stacking columns along the *a* axis. Neither stacking modes are favorable for a [2+2] photopolymerization reaction.

#### Comment

The crystal structures of a number of alkyl esters of  $\alpha$ -cyano-4-[2-(4-pyridyl)ethenyl]cinnamates have been analyzed in order to correlate their structures with their photoreactivity. For the methyl ester, photodimerization occurred between pyridyl-side olefinic groups of neighboring molecules. Oligomerization was observed on further photo-irradiation. For the ethyl and *n*-propyl homologs, however, dimerization occurred between the ester-side olefinic groups. Although the reaction terminated at the dimer formation for the *n*-propyl ester, the dimer of the ethyl ester gave a polymer on further irradiation (Maekawa, Kato, Saigo, Hasegawa & Ohashi, 1991; Hasegawa & Hashimoto, 1992; Hasegawa, 1995). This paper reports the structure of the *n*-butyl ester, (I). Recently, preliminary experiments on polymer formation have been performed and a variety of oligomeric compounds obtained. Although the products were investigated thoroughly, no high polymeric compounds were obtained.



There are two crystallographically independent molecules, A and B, both of which form infinite columns, A and B, along the a axis. The molecular structures

C3-C2-C7

<sup>†</sup> Alternative name: *n*-butyl 2-cyano-3-{4-[2-(4-pyridyl)ethenyl]phenyl}propenoate.



Fig. 1. ORTEPII (Johnson, 1976) drawing of the molecular structures and numbering systems of molecules A and B showing 50% probability displacement ellipsoids.



Fig. 2. Stacking patterns of columns A and B. The C—C distances (Å) between the double bonds are given.

and numbering schemes are shown in Fig. 1. The packing patterns of the A and B columns are different from each other, as shown in Fig. 2, in which the intermolecular distances between the ethylenic double bonds are given. In column A, the dimerization between the ester-side olefinic groups would easily occur easily on irradiation, considering the threshold value of 4.2 Å for the reaction (Schmidt, 1964, 1971). However, the oligomerization or polymerization may be difficult since the ethylenic double bonds of neighboring molecules are far from the pyridyl-side olefinic group. In column B, the pyridyl-side double bond is disordered. The intermolecular distances between the double bonds are within the threshold value. One of the disordered conformations is parallel and favorable for [2+2] cyclo-

addition. This indicates that the cyclic dimer would be produced. Another possibility is that the trimer may be produced since the distance between the ester-side double bonds (4.28 Å) is very close to the threshold value. In either case, polymerization in the crystal of the *n*-butyl ester is not very likely, which is consistent with the preliminary result. The dimers or trimers produced would be converted to various oligomers via nontopochemical reactions.

## Experimental

The title compound was prepared as described previously (Ichimura & Watanabe, 1982) and recrystallized from a benzene solution.

## Data collection

Rigaku R-AXIS IICS diffractometer

3934 reflections with  $I > 2\sigma(I)$ 

Symmetry $1 - x, 1 - x$	codes: (i) $y, 2 - z;$ (iv	1 - x, 2 - y, (1) $2 - x, 1 - 1$	2 - z; y, 2 - z	( :.
The data	a collection (	anditions h		v

The data collection conditions by R-AXIS IICS are as follows: number of frames measured 19, oscillation range per frame  $10.0^{\circ}$ , exposure time per frame 45 min and crystal-to-detector distance 143.0 mm.

Table 1. Selected intermolecular distances (Å)

C35...C36<sup>iii</sup>

C35. · · C36<sup>iv</sup>

 $C36 \cdot \cdot \cdot C35^{iii}$ 

 $C36 \cdot \cdot \cdot C35^{iv}$ 

3.780 (3)

4.273 (3)

3.936 (3)

4 273 (3)

 $R_{\rm int} = 0.053$ 

 $\theta_{\rm max} = 27.70^{\circ}$ 

 $h = -10 \rightarrow 9$ 

 $k = -34 \rightarrow 35$ 

 $l = -10 \rightarrow 10$ 

 $(\Delta/\sigma)_{\rm max} = -0.001$  $\Delta\rho_{\rm max} = 0.160 \text{ e} \text{ Å}^{-3}$ 

 $\Delta \rho_{\rm min} = -0.184 \ {\rm e} \ {\rm \AA}^{-3}$ 

Scattering factors from

Extinction correction: none

International Tables for

Crystallography (Vol. C)

2 - z; (ii) -x, 2 - y, 1 - z; (iii)

3.925 (3)

5.798 (3)

3.925 (3)

5.798 (3)

Data collection: OSCILL in R-AXIS IICS Software (Rigaku Corporation, 1994). Cell refinement: SCALE in R-AXIS IICS Software. Data reduction: SCALE in R-AXIS IICS Software. Program(s) used to solve structure: SIR92 (Altomare et al., 1994) in TEXSAN (Molecular Structure Corporation, 1995). Program(s) used to refine structure: SHELXL93 (Sheldrick, 1993). Molecular graphics: ORTEPII (Johnson, 1976).

Supplementary data for this paper are available from the IUCr electronic archives (Reference: DE1061). Services for accessing these data are described at the back of the journal.

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## $\beta$ -D-3'-Deoxyadenosine (Cordycepin)

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(Received 4 November 1996; accepted 15 May 1997)

### Abstract

In the title compound,  $C_{10}H_{13}N_5O_3$ , the nucleoside adopts an *anti* conformation with C3'-*endo* sugar puckering. The overall molecular conformation is similar to that of both adenosine and 2'-deoxyadenosine.

## Comment

3'-Deoxynucleosides could function as possible inhibitors of RNA synthesis and, thus, may be useful in the treatment of cancer, as well as viral and other infections (Bazin & Chattopadhyaya, 1985; Rainny & Santi, 1983). As part of our program to study their molecular conformations (Karthe, Gautham, Kumar & Katti, 1997), we report here the structure of 3'-deoxyadenosine, (I).



An ORTEP92 (Vicković, 1994) view of the molecule with the atom-numbering scheme is shown in Fig. 1. The conformation of the base with respect to the ribose moiety is anti with the glycosidic torsion angle  $\chi_{CN}$ , O4'—C1'—N9—C4, equal to -172.5 (6)°. In  $\alpha$ -D-2'deoxyadenosine (Watson, Sutor & Tollin, 1965) and also in  $\beta$ -D-adenosine (Lai & Marsh, 1972), the conformation is anti. The sugar pucker in the present structure is C3'-endo, as observed in adenosine. The pseudorotation phase angle (P) and the maximum amplitude of puckering ( $\tau_{M}$ ) are 4.4 and 38.4°, respectively. In 2'-deoxyadenosine, the sugar pucker is C3'-exo. The torsion angles  $\varphi_{OO}$  (O5'—C5'—C4'—O4') and  $\varphi_{OC}$  (O5'—C5'—C4'—C3') are 57.3 (8) and 175.5 (6)°, respectively. The conformation about the C4'—C5' bond

Refinement

S = 1.000

C14...C7<sup>i</sup>

 $C15 \cdot \cdot \cdot C6^i$ 

C14...C15<sup>ii</sup>

C15...C14<sup>ii</sup>

Refinement on  $F^2$ 

7740 reflections

539 parameters

H atoms not refined

+ 0.1876*P*] where  $P = (F_o^2 + 2F_c^2)/3$ 

 $w = 1/[\sigma^2(F_o^2) + (0.0639P)^2$ 

 $\frac{R[F^2 > 2\sigma(F^2)]}{wR(F^2)} = 0.051$ 

Area detector scans

Absorption correction: none

15 309 measured reflections

7744 independent reflections