$\begin{array}{c} 02C101\\ 02C1C2\\ 01C1C2\\ 03C2C1\\ 03C2C3\\ C1C2C3\\ 04C3C2\\ \end{array}$	126.6 (3)	04-C3-C4	112.0 (3)
	117.4 (3)	C4-C3-C2	110.0 (3)
	116.0 (3)	O5-C4-O6	125.3 (3)
	113.2 (3)	O5-C4-C3	121.7 (3)
	107.4 (3)	O6-C4-C3	113.0 (3)
	108.4 (3)	N1-C5-C6	110.7 (3)
	111.6 (3)	N2-C6-C5	110.6 (3)
$\begin{array}{c} 04-C3-C4-05\\ C2C3C4-05\\ 04C3C4-06\\ C2C3C4-06\\ 02C1C2-03\\ 01C1C2-03\\ 02C1C2C3\\ \end{array}$	-23.5 (4) 101.1 (4) 156.5 (3) -78.8 (3) 167.4 (3) -13.3 (5) -73.5 (4)	01C1C2C3 04C3C203 C4C3C203 04C3C2C1 C4C3C2C1 N1C5C6N2	105.8 (4) 58.0 (3) -67.0 (3) -64.7 (3) 170.4 (2) -60.6 (5)

Table 3. Least-squares-planes data

Plane 1: C1, C2, C3, C4; equation: -0.375(30)x + 2.477(11)y + 14.815(6)z = 9.121(18). Plane 2: C1, C2, O1, O2, O3; equation: 6.589(4)x - 1.360(15)y + 6.723(22)z = 4.785(20). Plane 3: C3, C4, O4, O5, O6; equation: -6.851(3)x - 2.909(9)y - 1.343(32)z = 2.066(15).

Dihedral angles (°)

Plane 1/Plane 2	72.37 (24)	Plane 2/Plane 3	32.16 (12)
Plane 1/Plane 3	75.49 (25)		

Deviations of atoms (Å) from their least-squares planes

	Plane 1	Plane 2	Plane 3
C1	0.060 (2)	-0.012 (3)	
C2	-0.058 (2)	-0.109 (2)	
C3	-0.063 (2)		-0.199 (2)
C4	0.061 (2)		-0.032 (3)
01		-0.035(1)	
02		0.068 (2)	
O3		0.089 (2)	
04			0.158 (2)
05			-0.049(1)
06			0.123 (2)

H atoms, except H1, H2A and H2B, were obtained by difference Fourier syntheses and were refined isotropically. Atoms H1, H2A and H2B were located geometrically.

Data collection: CAD-4-PC (Enraf-Nonius, 1993). Cell refinement: CAD-4-PC. Data reduction: MolEN (Fair, 1990). Program(s) used to solve structure: SHELXS86 (Sheldrick, 1990). Program(s) used to refine structure: SHELXL93 (Sheldrick, 1993). Molecular graphics: ORTEPII (Johnson, 1976). Software used to prepare material for publication: SHELXL93.

The authors thank Dr M. Kepez for supplying the crystal.

Lists of structure factors, anisotropic displacement parameters, Hatom coordinates and complete geometry have been deposited with the IUCr (Reference: KA1121). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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3-Acetoxymethylpyrrole-2-carbaldehyde

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Abstract

In the title compound (the ester 2-formyl-3-pyrrolylmethyl acetate, $C_8H_9NO_3$), hydrogen bonding between the N—H function in one molecule and the ester carbonyl O atom in a molecule related to the first by the 2_1 screw axis leads to the formation of zigzag chains.

Comment

3-Acetoxymethylpyrrole-2-carbaldehyde, (1), was prepared as an intermediate in a new synthetic route to 3,8-didehydroheliotridin-5-one (McNab & Thornley, 1993). Because of interest in the crystal structures of 3substituted pyrrole-2-carbaldehyde derivatives [(2)–(4)] (Smith, Bobe, Minnetian, Hope & Yanuck, 1985), we undertook a determination of its structure at 150 K.



Ring bond lengths in (1) follow the same pattern as seen in compounds (2)–(4) (Smith *et al.*, 1985). Thus, the substitution at the 2 and 3 positions of the pyrrole ring in all four compounds is reflected in the C2–C3 bond being consistently longer than the C4–C5 bond. Similarly, the N1–C2 bond in (1) is significantly longer than that between N1 and C5, a trend also observed in (2)–(4); this can be rationalized by conjugative electron donation from the lone pair of the N atom to the aldehyde function.

The ring angles in (1) are very similar to those in the methyl analogue (2), being widest at N1 and C5. The angles to each substituent are widened in the direction of the other substituent, particularly at C2 where the C3—C2—C21 angle is $130.6(3)^{\circ}$.

The O atom (O21) of the aldehyde function is twisted so as to lie 0.167(5) Å from the plane of the pyrrole ring. The acetoxymethyl substituent (C6–C9) is essentially planar and its normal forms an angle of $76.42(11)^{\circ}$ with the normal to the pyrrole ring.



Fig. 1. A view of the molecule with atom-numbering scheme. Displacement ellipsoids enclose 50% probability surfaces.



Fig. 2. A packing diagram of the structure showing intermolecular hydrogen bonding between the N—H group and the carbonyl O atom of the ester function.

As can be seen from Fig. 2, the structure exhibits intermolecular hydrogen bonding between the N—H function and the carbonyl O atom of the ester function. These interactions link molecules related by the operation of the 2_1 screw axis into zigzag chains.

Experimental

The title compound was prepared by photolysis of 4-acetoxymethylpyridine N-oxide in aqueous copper(II) sulfate solution. Crystals could be obtained by slow evaporation from either ethanol or toluene.

Crystal data

C ₈ H ₉ NO ₃	Mo $K\alpha$ radiation
$M_r = 167.16$	$\lambda = 0.71073 \text{ Å}$
Monoclinic	Cell parameters from 24
$P2_1/n$	reflections
a = 4.463(3) Å	$\theta = 13 - 16^{\circ}$
b = 11.546(5) Å	$\mu = 0.107 \text{ mm}^{-1}$
c = 15.697(5) Å	T = 150.0 (2) K
$\beta = 95.75(5)^{\circ}$	Irregular block
$V = 804.8(7) \text{ Å}^3$	$0.43 \times 0.27 \times 0.19$ mm
Z = 4	Colourless
$D_r = 1.380 \text{ Mg m}^{-3}$	

 $R_{\rm int} = 0.0402$ $\theta_{\rm max} = 25.01^{\circ}$

 $h = -5 \rightarrow 5$ $k = -2 \rightarrow 13$

 $l = -11 \rightarrow 18$

3 standard reflections frequency: 60 min

intensity decay: 3%

Data collection

Stoe Stadi-4 four-circle diffractometer ω -2 θ scans using on-line profile fitting (Clegg, 1981) Absorption correction: none 1492 measured reflections 1404 independent reflections 1015 observed reflections $[I > 2\sigma(I)]$

Refinement

N1

C2 C3

 $\Delta \rho_{\rm max} = 0.23 \ {\rm e} \ {\rm \AA}^{-3}$ Refinement on F^2 $\Delta \rho_{\rm min} = -0.27 \ {\rm e} \ {\rm \AA}^{-3}$ $R[F^2 > 2\sigma(F^2)] = 0.0538$ $wR(F^2) = 0.1703$ Extinction correction: S = 0.983SHELXL93 (Sheldrick, 1392 reflections 1993) 146 parameters Extinction coefficient: All H-atom parameters 0.021(6)Atomic scattering factors refined $w = 1/[\sigma^2(F_o^2) + (0.0611P)^2]$ from International Tables + 1.37*P*] for Crystallography (1992, where $P = (F_o^2 + 2F_c^2)/3$ Vol. C. Tables 4.2.6.8 and $(\Delta/\sigma)_{\rm max} = -0.03$ 6.1.1.4)

 Table 1. Fractional atomic coordinates and equivalent isotropic displacement parameters (Å²)

$U_{\rm eq} = (1/3) \sum_i \sum_j U_{ij} a_i^* a_i^* \mathbf{a}_i \cdot \mathbf{a}_j.$

x	у	Z	U_{eq}
0.6335 (6)	-0.0829 (2)	0.7734 (2)	0.0267 (6)
0.4518 (7)	0.0135 (3)	0.7640 (2)	0.0244 (7)
0.3078 (6)	0.0242 (3)	0.8381 (2)	0.0238 (7)

C₈H₉NO₃

C4	0.4070(7)	-0.0675 (3)	0.8923 (2)	0.0284 (7)
C5	0.6045 (7)	-0.1319 (3)	0.8503 (2)	0.0295 (7)
C6	0.0956 (7)	0.1186(3)	0.8591 (2)	0.0272 (7)
07	0.2367 (5)	0.1971 (2)	0.92455 (13)	0.0284 (6)
C8	0.4232 (7)	0.2779 (2)	0.8981 (2)	0.0254 (7)
08	0.4685 (5)	0.2888 (2)	0.82381 (13)	0.0305 (6)
C9	0.5695 (10)	0.3479 (3)	0.9704 (2)	0.0393 (9)
C21	0.4430 (8)	0.0862 (3)	0.6898 (2)	0.0318 (8)
O21	0.6045 (6)	0.0738 (2)	0.63177 (15)	0.0456 (7)

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N1C5	1.351 (4)	C4—C5	1.370 (5)
N1C2	1.377 (4)	C6—O7	1.464 (4)
C2C3	1.389 (4)	O7C8	1.344 (4)
C2C21	1.433 (4)	C808	1.209 (4)
C3—C4	1.402 (4)	C8—C9	1.490 (5)
C3—C6	1.502 (4)	C21O21	1.225 (4)
C5N1C2	109.0 (3)	N1-C5-C4	109.1 (3)
N1-C2-C3	107.4 (3)	O7C6C3	111.5 (3)
N1-C2-C21	121.9 (3)	C8-07-C6	116.7 (2)
C3-C2-C21	130.6 (3)	O8C8O7	122.6 (3)
C2C3C4	107.2 (3)	08	125.2 (3)
C2-C3-C6	127.1 (3)	07	112.2 (3)
C4—C3—C6	125.7 (3)	O21-C21-C2	124.5 (3)
C5-C4-C3	107.3 (3)		
C2-C3-C6-07	-107.1 (3)	C6-07-C8-C9	-176.3 (3)
C4-C3-C6-07	70.0 (4)	N1-C2-C21-O21	-4.4 (5)
C3-C6-07-C8	79.4 (3)	C3-C2-C21-O21	172.3 (3)
C6-07-C8-08	2.1 (4)		

Data collection: *DIF*4 (Stoe & Cie, 1988*a*). Cell refinement: *DIF*4. Data reduction: *REDU*4 (Stoe & Cie, 1988*b*). Program(s) used to solve structure: *SHELXS*86 (Sheldrick, 1990). Program(s) used to refine structure: *SHELXL*93 (Sheldrick, 1993). Molecular graphics: *SHELXTL/PC* (Sheldrick, 1992).

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Lists of structure factors, anisotropic displacement parameters, Hatom coordinates and complete geometry have been deposited with the IUCr (Reference: PT1024). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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Dimethyl [α -(Benzylamino)-*p*-chlorobenzyl]phosphonate

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Abstract

The title compound, $C_{16}H_{19}CINO_3P$, is one of a series of α -substituted aminomethyl phosphonates. There are two independent molecules in the asymmetric unit and in each the P atom adopts a tetrahedral configuration. The P=O double-bond length and the mean P-O single-bond length are 1.425 (5) and 1.560 (5) Å for molecule (1) and 1.417 (6) and 1.562 (5) Å for molecule (2), respectively. The P=O double-bond lengths in both molecules are shorter than those in 1,3,2-oxazaphospholidine derivatives and caged bicyclic phosphorus compounds.

Comment

 α -Substituted aminomethyl phosphonates have been investigated (Li, Wang, Zhang, Miao, Liu, Cao & Guo, 1988; Li, Wang, Han, Liu, Cao, Jiang, Miao & Liu, 1988) because of their potential biological activity. The title compound, (I), was synthesized and its crystal structure is presented here.



The asymmetric unit contains two independent molecules. The P atom adopts a distorted tetrahedral configuration; the bond angles around the P atom are in the range $100.9(2)-116.1(3)^{\circ}$ for molecule (1) and $97.8(3)-119.3(3)^{\circ}$ for molecule (2). The mean P—O