

C25	0.1804 (2)	1.04731 (15)	0.6204 (2)	0.0416 (6)
C26	0.1482 (2)	1.02818 (13)	0.4750 (2)	0.0339 (5)
C31	0.2198 (2)	1.07360 (12)	0.1381 (2)	0.0249 (4)
C32	0.2996 (2)	1.13061 (13)	0.2166 (2)	0.0299 (5)
C33	0.3152 (2)	1.20978 (14)	0.1616 (2)	0.0377 (5)
C34	0.2521 (2)	1.23238 (14)	0.0263 (3)	0.0405 (6)
C35	0.1743 (2)	1.17634 (14)	-0.0540 (2)	0.0395 (6)
C36	0.1574 (2)	1.09760 (14)	0.0019 (2)	0.0331 (5)

Table 2. Selected geometric parameters (Å, °)

P1—C21	1.828 (2)	C11—C16	1.396 (3)
P1—C11	1.846 (2)	C11—C12	1.422 (3)
P1—C31	1.851 (2)	C12—C13	1.400 (3)
O1—C3	1.211 (2)	C13—C14	1.381 (3)
C1—C2	1.511 (3)	C14—C15	1.377 (3)
C2—C3	1.516 (3)	C15—C16	1.387 (3)
C3—C12	1.492 (3)		
C21—P1—C11	104.37 (9)	C13—C12—C3	120.2 (2)
C21—P1—C31	99.69 (9)	C11—C12—C3	120.2 (2)
C11—P1—C31	102.24 (9)	C14—C13—C12	121.3 (2)
C1—C2—C3	113.0 (2)	C15—C14—C13	119.7 (2)
O1—C3—C12	120.0 (2)	C14—C15—C16	120.0 (2)
O1—C3—C2	121.3 (2)	C15—C16—C11	122.1 (2)
C12—C3—C2	118.8 (2)	C22—C21—P1	125.4 (2)
C16—C11—C12	117.4 (2)	C26—C21—P1	116.4 (2)
C16—C11—P1	121.43 (15)	C36—C31—P1	115.8 (2)
C12—C11—P1	120.70 (14)	C32—C31—P1	125.6 (2)
C13—C12—C11	119.5 (2)		
O1—C3—C12—C11	-11.2 (2)	C1—C2—C3—O1	0.3 (2)

Data collection: XSCANS (Siemens, 1993). Cell refinement: XSCANS. Data reduction: XSCANS. Program(s) used to solve structure: SHELXS86 (Sheldrick, 1990). Program(s) used to refine structure: SHELXL93 (Sheldrick, 1993). Molecular graphics: SHELXTL-Plus (Sheldrick, 1987). Software used to prepare material for publication: SHELXL93.

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Lists of structure factors, anisotropic displacement parameters, H-atom coordinates and complete geometry have been deposited with the IUCr (Reference: FG1009). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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2,6-Diphenyl-3-vinylpyridine

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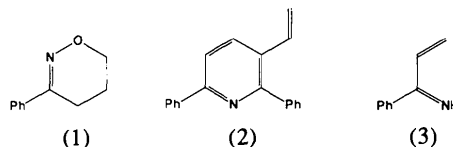
(Received 29 June 1994; accepted 19 September 1994)

Abstract

In the structure of C₁₉H₁₅N, steric interactions force all three substituents (one vinyl and two phenyl groups) out of coplanarity with the central pyridine ring. Individual torsion angles around the relevant bonds are in the ranges 33.2 (3)–36.1 (3), 38.9 (3)–41.8 (3) and 29.6 (3)–31.1 (3)°.

Comment

In connection with an investigation of the chemistry of 5,6-dihydro-4H-1,2-oxazines (Ellames, Hewkin, Jackson, Smith & Standen, 1989), we treated 3-phenyl-5,6-dihydro-4H-1,2-oxazine, (1), with potassium *t*-amyloxide (6 equivalents) in *t*-amyl alcohol at reflux for 4 h. Flash chromatography allowed the isolation of the title compound (2) (13% yield) and the structure was determined by X-ray crystallography. This structure is consistent with the spectroscopic data obtained.



We believe that the title compound arises as a result of a Diels–Alder cycloaddition reaction between the imine (3) [which we have already postulated as an intermediate in the base-induced reactions of oxazine (1)] and 1-phenylbuta-1,3-diene, also derived from oxazine (1) by base-induced double-bond migration and a retro-Diels–Alder reaction. A final air oxidation would then lead to the aromatic title compound (2).

Although a number of copper and nickel complexes of 2-vinylpyridine have been structurally characterized (Engelhardt, Healy, Kildea & White, 1989; Hoberg, Guhl & Betz, 1990; Munakata, Kitagawa, Simono, Emori & Masuda, 1987), this appears to be the first report of the crystal structure of a 3-vinylpyridine, or indeed of any vinylpyridine not complexed to a metal. The three aromatic rings are essentially planar (r.m.s. deviations of 0.004, 0.003 and 0.004 Å), but the two phenyl and one vinyl substituent are twisted markedly

out of the plane of the central pyridine ring by 40.4, 30.4 and 34.6°, respectively. This entails a loss of conjugation, but is necessitated by the steric interactions of the H atoms, as is commonly observed for connected aromatic rings.

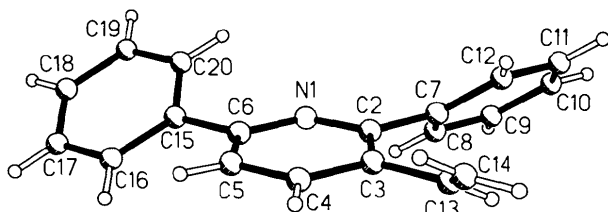


Fig. 1. Molecular structure of the title compound with atom labels.

Experimental

A suitable crystal was obtained by evaporation of the appropriate fractions following chromatography.

Crystal data

C₁₉H₁₅N
M_r = 257.32
 Monoclinic
*P*2₁/*c*
a = 11.8386 (9) Å
b = 15.3158 (12) Å
c = 7.7899 (6) Å
 β = 95.970 (8)°
V = 1404.8 (2) Å³
Z = 4
D_x = 1.217 Mg m⁻³

Data collection

Stoe Siemens diffractometer
 ω/θ scans with on-line
 profile fitting (Clegg,
 1981)
 Absorption correction:
 none
 2277 measured reflections
 2264 independent reflections
 2014 observed reflections
 [*I* > 2σ(*I*)]

Refinement

Refinement on *F*²
R(*F*) = 0.0588
wR(*F*²) = 0.1739
S = 1.059
 2260 reflections
 182 parameters
 H atoms riding with *U*(H) =
 1.2*U*_{eq}(C)
w = 1/[σ²(*F*_o²) + (0.1134*P*)²
 + 0.2574*P*]
 where *P* = (*F*_o² + 2*F*_c²)/3
 (Δ/σ)_{max} < 0.001

Cu *K*α radiation
 λ = 1.5418 Å
 Cell parameters from 32
 reflections
 θ = 15.0–20.0°
 μ = 0.538 mm⁻¹
T = 295 (2) K
 Block
 0.50 × 0.40 × 0.40 mm
 Colourless

*R*_{int} = 0.0617
 θ_{max} = 65.02°
h = -13 → 13
k = -17 → 18
l = -9 → 9
 3 standard reflections
 frequency: 60 min
 intensity decay: 1%

Δρ_{max} = 0.26 e Å⁻³
 Δρ_{min} = -0.21 e Å⁻³
 Extinction correction:
 SHELXL93 (Sheldrick,
 1993)
 Extinction coefficient:
 0.0073 (12)
 Atomic scattering factors
 from *International Tables*
 for *Crystallography* (1992,
 Vol. C, Tables 4.2.6.8 and
 6.1.1.4)

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

$$U_{\text{eq}} = (1/3)\sum_i \sum_j U_{ij} a_i^* a_j^*$$

	<i>x</i>	<i>y</i>	<i>z</i>	<i>U</i> _{eq}
N1	0.55484 (12)	0.35169 (9)	0.2542 (2)	0.0444 (4)
C2	0.65647 (14)	0.32838 (11)	0.2074 (2)	0.0417 (4)
C3	0.6905 (2)	0.24060 (11)	0.1973 (2)	0.0442 (5)
C4	0.6124 (2)	0.17766 (12)	0.2367 (2)	0.0507 (5)
C5	0.5090 (2)	0.20144 (12)	0.2863 (2)	0.0510 (5)
C6	0.48146 (15)	0.28963 (11)	0.2928 (2)	0.0443 (5)
C7	0.72857 (15)	0.40341 (11)	0.1641 (2)	0.0430 (5)
C8	0.6770 (2)	0.47393 (12)	0.0740 (2)	0.0517 (5)
C9	0.7410 (2)	0.54497 (13)	0.0319 (3)	0.0598 (6)
C10	0.8552 (2)	0.54777 (13)	0.0804 (3)	0.0628 (6)
C11	0.9076 (2)	0.47892 (14)	0.1716 (3)	0.0590 (6)
C12	0.8444 (2)	0.40763 (12)	0.2128 (2)	0.0500 (5)
C13	0.8002 (2)	0.21386 (12)	0.1406 (2)	0.0519 (5)
C14	0.8588 (2)	0.14516 (15)	0.1994 (3)	0.0695 (7)
C15	0.37090 (15)	0.32083 (12)	0.3429 (2)	0.0460 (5)
C16	0.2723 (2)	0.27216 (14)	0.3082 (3)	0.0575 (5)
C17	0.1699 (2)	0.3031 (2)	0.3533 (3)	0.0699 (7)
C18	0.1646 (2)	0.3823 (2)	0.4353 (3)	0.0712 (7)
C19	0.2611 (2)	0.4313 (2)	0.4708 (3)	0.0661 (6)
C20	0.3637 (2)	0.40137 (13)	0.4238 (3)	0.0551 (5)

Table 2. Selected geometric parameters (Å, °)

N1—C2	1.341 (2)	C8—C9	1.385 (3)
N1—C6	1.343 (2)	C9—C10	1.366 (3)
C2—C3	1.408 (2)	C10—C11	1.382 (3)
C2—C7	1.491 (2)	C11—C12	1.381 (3)
C3—C4	1.392 (3)	C13—C14	1.316 (3)
C3—C13	1.472 (3)	C15—C16	1.388 (3)
C4—C5	1.371 (3)	C15—C20	1.392 (3)
C5—C6	1.392 (3)	C16—C17	1.381 (3)
C6—C15	1.483 (2)	C17—C18	1.376 (3)
C7—C12	1.386 (3)	C18—C19	1.371 (3)
C7—C8	1.394 (3)	C19—C20	1.383 (3)
C2—N1—C6	119.48 (15)	N1—C6—C15	116.1 (2)
N1—C2—C3	122.6 (2)	C5—C6—C15	122.6 (2)
N1—C2—C7	114.03 (14)	C12—C7—C8	118.1 (2)
C3—C2—C7	123.3 (2)	C12—C7—C2	123.2 (2)
C4—C3—C2	116.6 (2)	C8—C7—C2	118.7 (2)
C4—C3—C13	120.0 (2)	C14—C13—C3	124.6 (2)
C2—C3—C13	123.3 (2)	C16—C15—C20	118.3 (2)
C5—C4—C3	120.8 (2)	C16—C15—C6	121.5 (2)
C4—C5—C6	119.2 (2)	C20—C15—C6	120.1 (2)
N1—C6—C5	121.3 (2)		
Phenyl C—C—C	119.8 (2)–121.2 (2)		
C14—C13—C3—C2	-36.1 (3)	C3—C2—C7—C12	-41.8 (3)
C14—C13—C3—C2	146.8 (3)	N1—C6—C15—C16	149.1 (3)
N1—C2—C7—C8	-38.9 (3)	N1—C6—C15—C20	-29.6 (3)
N1—C2—C7—C12	139.4 (3)	C5—C6—C15—C16	-31.1 (3)
C3—C2—C7—C8	139.9 (3)	C5—C6—C15—C20	150.1 (3)

The data set before averaging consisted of a complete unique set of reflections (*k*, *l* ≥ 0), together with a few Friedel opposites, which are responsible for the index limits given.

Data collection: *DIF4* (Stoe & Cie, 1988). Cell refinement: *DIF4*. Data reduction: local programs. Program(s) used to solve structure: *SHELXTL/PC* (Sheldrick, 1990). Program(s) used to refine structure: *SHELXL93* (Sheldrick, 1993). Molecular graphics: *SHELXTL/PC*. Software used to prepare material for publication: *SHELXL93* and local programs.

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Lists of structure factors, anisotropic displacement parameters, H-atom coordinates, complete geometry and torsion angles have been deposited with the IUCr (Reference: MU1138). Copies may be obtained through The Managing Editor, International Union of Crystallography, 5 Abbey Square, Chester CH1 2HU, England.

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Acta Cryst. (1995). **C51**, 527–529

2,2-Dimethyl-5-(2-methyl-4-phenyl-3,4-diaza-2-butenylidene)-1,3-dioxane-4,6-dione

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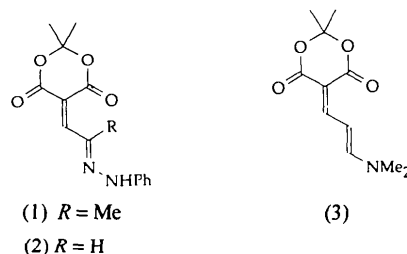
(Received 13 June 1994; accepted 15 July 1994)

Abstract

The molecular conformation of the title compound, $C_{15}H_{16}N_2O_4$, is described in terms of three planar sections which are mutually twisted. The main intermolecular contacts are $N-H\cdots O$ hydrogen bonds where the $O\cdots N$ separation is 3.063 (3) Å.

Comment

The study of the structure of the title compound, (1), was undertaken to determine whether there were conformational reasons for its ready cyclization to give a 3-oxopyridazinecarboxylic acid upon thermolysis in solution, while the corresponding unsubstituted compound (2) is recovered unchanged under similar conditions (McNab & Stobie, 1982). The structure of (1) is also related to that of the propenylidene Meldrum's acid derivative (3), which we have described recently (Blake, McNab & Monahan, 1991).



The conjugated system from C5 to N10 in (1) is exclusively *trans*, as found for (3), and so the cyclization reaction cannot be explained by a ground-state proximity of N10 and C4.

A feature of the structure of (3) is that electron delocalization occurs along the propenyl chain such that the lengths of all the C—C and C=C bonds are comparable [1.385 (8)–1.409 (8) Å]. In the case of (1), the system is much more localized [C=C 1.355 (3), C—C 1.439 (3) Å], partly due to poorer electron donation from the hydrazone N atom N10 compared with the terminal amino group in (3), and also partly due to the non-planarity of the system (see below). This reduced electron donation is also reflected in the lengths of the C4—C5 and C5—C6 bonds [both 1.469 (3) Å], which are substantially longer than the corresponding values for (3) [1.449 (8), 1.417 (8) Å].

The presence of the C8M methyl substituent in (1) has a dramatic effect on the angles subtended at C8 and C5. Non-bonded contacts between the C8M methyl group and O4 lead to widening of the C7—C8—C8M and C8M—C8—N9 angles to 125.7 (2) and 123.7 (2)°, respectively, with a concomitant reduction in the C7—C8—N9 angle to 110.5 (2)°. The C4—C5—C7 angle is widened to 126.0 (2)°, compared with an angle of 120.9 (5)° in (3). This distortion is almost entirely at the expense of the endocyclic C4—C5—C6 angle [116.5 (2)°, compared with 121.7 (5)° in (3)]. A similar trend at C5 has been noted for the corresponding angles in the dimethylaminomethylene Meldrum's acid, which has analogous steric constraints (Blake, McNab & Monahan, 1991). In (1), the two exocyclic N—C—C angles involving the phenyl group differ significantly, possibly because of repulsion between the *ortho* H6P atom and the lone pair on N9.

The molecular conformation of (1) can be described in terms of three planar sections: the phenyl ring, the five-atom chain C5—C7—C8—N9—N10 with its C8M substituent, and four atoms (O1, O3, C4 and C6) of the Meldrum's acid ring. The angle between the normals to the phenyl ring and the chain is 10.2 (2)° and the angle between the normals to the chain and the plane of the Meldrum's acid ring is 20.5 (2)°. In contrast, there is a much greater degree of coplanarity between the chain and the Meldrum's acid ring in (3). The mean deviation from the plane of the chain is 0.058 Å, while the four atoms defining the Meldrum's acid plane