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1-Vinyl-1H-indole-3-carbaldehyde

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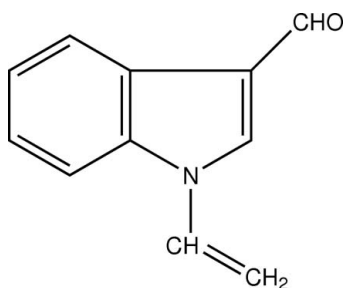
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 Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.002$ Å; R factor = 0.045; wR factor = 0.127; data-to-parameter ratio = 17.6.

In the title compound, $\text{C}_{11}\text{H}_9\text{NO}$, the C and O atoms of the attached carbaldehyde group deviate by just 0.052 (2) and 0.076 (1) Å, respectively, from the mean plane of the indole ring system. In addition to van der Waals forces, the molecular packing is stabilized by C—H \cdots O hydrogen bonds, which form a $C(7)$ chain motif, and π – π interactions (centroid–centroid distance 3.637 Å) between the pyrrole and benzene rings of the indole ring system.

Related literature

For related literature, see: Padwa *et al.* (1999); Mathiesen *et al.* (2005); Grinev *et al.* (1984); Gadaginamath & Patil (1999); Rodriguez *et al.* (1985); Karthick *et al.* (2005); Selvanayagam *et al.* (2005); Sonar *et al.* (2005). For bond-length data, see: Allen *et al.* (1987).



Experimental

Crystal data

 $\text{C}_{11}\text{H}_9\text{NO}$
 $M_r = 171.19$
 Monoclinic, $P2_1/n$
 $a = 8.3200$ (5) Å

 $b = 8.1490$ (5) Å
 $c = 13.1620$ (7) Å
 $\beta = 99.952$ (1)°
 $V = 878.95$ (9) Å³
 $Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 0.08$ mm⁻¹
 $T = 293$ (2) K
 $0.24 \times 0.22 \times 0.20$ mm

Data collection

 Bruker SMART APEX CCD area-detector diffractometer
 Absorption correction: none
 9730 measured reflections

 2072 independent reflections
 1823 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.020$

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.044$
 $wR(F^2) = 0.127$
 $S = 1.05$
 2072 reflections

 118 parameters
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.21$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.19$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{C9}-\text{H9}\cdots\text{O1}^i$	0.93	2.51	3.390 (2)	159

 Symmetry code: (i) $x - \frac{1}{2}, -y + \frac{1}{2}, z + \frac{1}{2}$.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT* (Bruker, 2001); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *ORTEP-3* (Farrugia, 1997) and *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXL97* and *PARST* (Nardelli, 1995).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BT2714).

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1-Vinyl-1*H*-indole-3-carbaldehyde

S. Selvanayagam, B. Sridhar, K. Ravikumar, S. Kathiravan and R. Raghunathan

S1. Comment

Indoles and their derivatives have been interest for many years, since large number of natural products contain indole systems and they are found in a number of pharmaceutical products, fragrances and dyes (Padwa *et al.*, 1999). Indole derivatives are identified as interfering with a G protein-independent signaling pathway of the CRTH2 receptor (Mathiesen *et al.*, 2005). These derivatives possess antidepressant (Grinev *et al.*, 1984), anti-microbial (Gadaginamath & Patil, 1999) and anti-inflammatory (Rodriguez *et al.*, 1985) activities. In view of its importance, we have undertaken the single-crystal X-ray diffraction study and report here its results.

The X-ray study confirmed the molecular structure and atomic connectivity for (I), as illustrated in Fig. 1. The geometry of the indole ring system is comparable to those reported for other indole derivatives (Karthick *et al.*, 2005; Selvanayagam *et al.*, 2005; Sonar *et al.*, 2005). The bond length of C9—C10 [1.284 (2) Å] confirms the double bond character (Allen *et al.*, 1987). The sum of the angles at N1 of the indole ring (360°) is in accordance with sp^2 hybridization.

The indole ring is planar with a maximum deviation of 0.017 (1) Å for atom C8. The carbaldehyde group atoms C11 and O1 deviate 0.052 (2) and 0.076 (1) Å, respectively from the best plane of the indole ring.

In addition to the van der Waals forces, the molecular packing is stabilized by intermolecular C—H \cdots O hydrogen bond (Table 2). Atom H9 of C9 forms a intermolecular hydrogen bond with oxygen atom O1 forming a C(7) chain motif of C—H \cdots O hydrogen bond along the diagonal of *ac* plane (Fig. 2). In addition to this a weak $\pi\cdots\pi$ interaction between the pyrrole ring (N1/C1/C6—C8) at (*x,y,z*) and benzene ring (C1—C6) at (1 -*x*, -*y*), -*z*) stabilizes the molecular packing. The centroid-to-centroid distance is 3.637 Å.

S2. Experimental

A mixture of *N*-vinylindole (0.05 mol) and DMF (0.15 mol) was stirred with POCl₃ (32.3 ml). The reaction mixture was poured into ice water (300 ml) and stirred for 30 minutes at less than 10° C. The precipitated solid was collected by filtration and washed well water (100 ml). In order to get the diffraction quality crystals, the compound was recrystallized from ethyl acetate.

S3. Refinement

The H atoms were positioned geometrically with C—H distances of 0.93 Å and were included in the refinement in the riding motion approximation with $U_{iso} = 1.2U_{eq}(C)$.

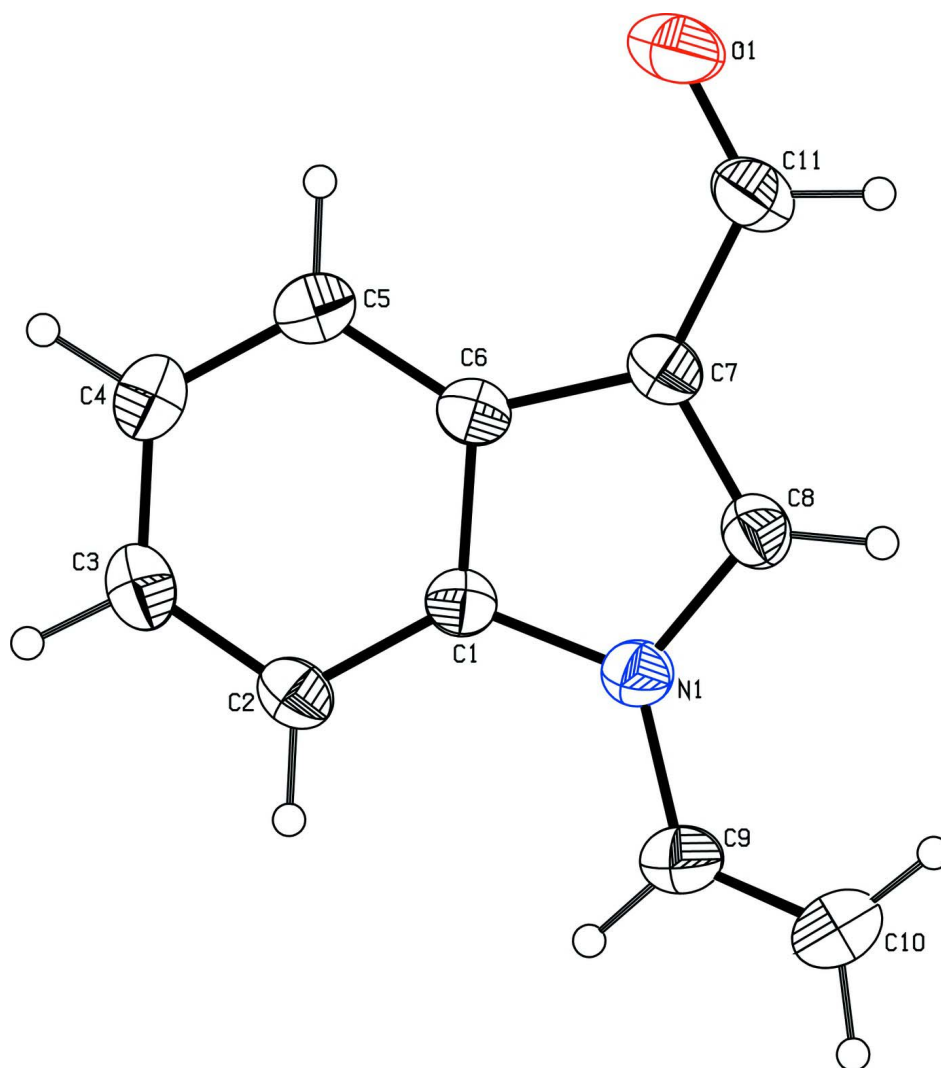
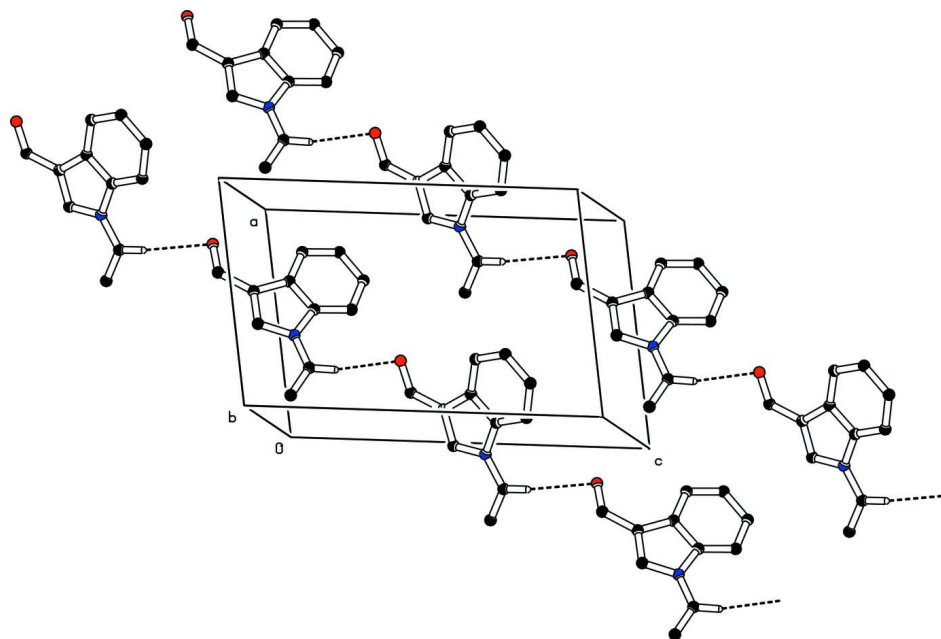


Figure 1

The structure and atom-numbering scheme for the title compound; displacement ellipsoids are drawn at the 30% probability level and H atoms are shown as small spheres of arbitrary radius.

**Figure 2**

Molecular packing of the title compound viewed down the *b* axis; H-bonds are shown as dashed lines. For clarity, H atoms, not involved in hydrogen bonds, have been omitted.

1-Vinyl-1H-indole-3-carbaldehyde

Crystal data

$C_{11}H_9NO$

$M_r = 171.19$

Monoclinic, $P2_1/n$

Hall symbol: -P 2yn

$a = 8.3200 (5) \text{ \AA}$

$b = 8.1490 (5) \text{ \AA}$

$c = 13.1620 (7) \text{ \AA}$

$\beta = 99.952 (1)^\circ$

$V = 878.95 (9) \text{ \AA}^3$

$Z = 4$

$F(000) = 360$

$D_x = 1.294 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$

Cell parameters from 4554 reflections

$\theta = 2.1\text{--}23.6^\circ$

$\mu = 0.08 \text{ mm}^{-1}$

$T = 293 \text{ K}$

Block, colourless

$0.24 \times 0.22 \times 0.20 \text{ mm}$

Data collection

Bruker SMART APEX CCD area-detector
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

ω scans

9730 measured reflections

2072 independent reflections

1823 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.020$

$\theta_{\text{max}} = 28.0^\circ$, $\theta_{\text{min}} = 3.0^\circ$

$h = -10 \rightarrow 10$

$k = -10 \rightarrow 10$

$l = -17 \rightarrow 16$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.044$

$wR(F^2) = 0.127$

$S = 1.05$

2072 reflections

118 parameters

0 restraints

Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier map
 Hydrogen site location: inferred from neighbouring sites
 H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0665P)^2 + 0.1375P]$$

where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.21 \text{ e } \text{Å}^{-3}$
 $\Delta\rho_{\min} = -0.19 \text{ e } \text{Å}^{-3}$

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) *etc.* and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (Å²)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	0.81283 (15)	0.20464 (18)	-0.13362 (9)	0.0826 (4)
N1	0.41791 (12)	0.24005 (13)	0.07004 (7)	0.0469 (3)
C1	0.54641 (14)	0.14623 (13)	0.12226 (8)	0.0424 (3)
C2	0.56155 (16)	0.06833 (16)	0.21713 (9)	0.0505 (3)
H2	0.4800	0.0748	0.2572	0.061*
C3	0.70276 (18)	-0.01899 (17)	0.24918 (10)	0.0581 (3)
H3	0.7166	-0.0736	0.3121	0.070*
C4	0.82544 (17)	-0.02764 (17)	0.18966 (11)	0.0594 (4)
H4	0.9197	-0.0870	0.2139	0.071*
C5	0.80997 (15)	0.04980 (16)	0.09564 (10)	0.0527 (3)
H5	0.8925	0.0434	0.0563	0.063*
C6	0.66781 (14)	0.13815 (13)	0.06061 (8)	0.0431 (3)
C7	0.60762 (15)	0.23003 (15)	-0.03148 (9)	0.0480 (3)
C8	0.45684 (16)	0.28725 (16)	-0.02188 (9)	0.0503 (3)
H8	0.3901	0.3497	-0.0712	0.060*
C9	0.27187 (17)	0.27412 (19)	0.10596 (12)	0.0615 (4)
H9	0.2701	0.2521	0.1751	0.074*
C10	0.14066 (19)	0.3327 (2)	0.05257 (15)	0.0779 (5)
H10A	0.1365	0.3567	-0.0169	0.094*
H10B	0.0498	0.3511	0.0834	0.094*
C11	0.68162 (19)	0.2565 (2)	-0.12120 (11)	0.0621 (4)
H11	0.6233	0.3191	-0.1742	0.075*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0720 (7)	0.1185 (10)	0.0649 (7)	0.0021 (6)	0.0336 (6)	0.0075 (6)
N1	0.0474 (5)	0.0518 (6)	0.0428 (5)	0.0032 (4)	0.0110 (4)	-0.0026 (4)
C1	0.0457 (6)	0.0418 (5)	0.0396 (5)	-0.0022 (4)	0.0075 (4)	-0.0069 (4)
C2	0.0595 (7)	0.0521 (7)	0.0417 (6)	-0.0035 (5)	0.0136 (5)	-0.0023 (5)
C3	0.0707 (8)	0.0549 (7)	0.0464 (6)	0.0007 (6)	0.0035 (6)	0.0051 (5)

C4	0.0562 (7)	0.0544 (7)	0.0641 (8)	0.0080 (6)	0.0011 (6)	0.0007 (6)
C5	0.0475 (6)	0.0527 (7)	0.0591 (7)	0.0000 (5)	0.0129 (5)	-0.0070 (5)
C6	0.0472 (6)	0.0418 (6)	0.0411 (5)	-0.0053 (4)	0.0102 (4)	-0.0072 (4)
C7	0.0543 (7)	0.0486 (6)	0.0425 (6)	-0.0046 (5)	0.0124 (5)	-0.0023 (5)
C8	0.0566 (7)	0.0513 (7)	0.0427 (6)	0.0017 (5)	0.0079 (5)	0.0021 (5)
C9	0.0564 (8)	0.0736 (9)	0.0586 (8)	0.0089 (6)	0.0209 (6)	-0.0013 (6)
C10	0.0580 (9)	0.0892 (12)	0.0894 (12)	0.0170 (8)	0.0205 (8)	0.0036 (9)
C11	0.0670 (8)	0.0743 (9)	0.0481 (7)	-0.0059 (7)	0.0185 (6)	0.0041 (6)

Geometric parameters (Å, °)

O1—C11	1.2079 (19)	C5—C6	1.3933 (17)
N1—C8	1.3610 (16)	C5—H5	0.9300
N1—C1	1.3949 (15)	C6—C7	1.4390 (17)
N1—C9	1.4055 (16)	C7—C8	1.3645 (18)
C1—C2	1.3869 (16)	C7—C11	1.4393 (18)
C1—C6	1.4023 (16)	C8—H8	0.9300
C2—C3	1.3760 (19)	C9—C10	1.284 (2)
C2—H2	0.9300	C9—H9	0.9300
C3—C4	1.392 (2)	C10—H10A	0.9300
C3—H3	0.9300	C10—H10B	0.9300
C4—C5	1.3751 (19)	C11—H11	0.9300
C4—H4	0.9300		
C8—N1—C1	108.24 (10)	C5—C6—C1	119.20 (11)
C8—N1—C9	126.55 (12)	C5—C6—C7	134.39 (11)
C1—N1—C9	125.17 (11)	C1—C6—C7	106.40 (10)
C2—C1—N1	129.57 (11)	C8—C7—C6	106.95 (10)
C2—C1—C6	122.48 (11)	C8—C7—C11	123.72 (13)
N1—C1—C6	107.94 (10)	C6—C7—C11	129.29 (12)
C3—C2—C1	116.91 (12)	N1—C8—C7	110.46 (11)
C3—C2—H2	121.5	N1—C8—H8	124.8
C1—C2—H2	121.5	C7—C8—H8	124.8
C2—C3—C4	121.60 (12)	C10—C9—N1	126.30 (15)
C2—C3—H3	119.2	C10—C9—H9	116.8
C4—C3—H3	119.2	N1—C9—H9	116.8
C5—C4—C3	121.31 (12)	C9—C10—H10A	120.0
C5—C4—H4	119.3	C9—C10—H10B	120.0
C3—C4—H4	119.3	H10A—C10—H10B	120.0
C4—C5—C6	118.49 (12)	O1—C11—C7	125.68 (15)
C4—C5—H5	120.8	O1—C11—H11	117.2
C6—C5—H5	120.8	C7—C11—H11	117.2
C8—N1—C1—C2	-178.34 (12)	N1—C1—C6—C7	-0.32 (12)
C9—N1—C1—C2	-0.3 (2)	C5—C6—C7—C8	178.96 (13)
C8—N1—C1—C6	0.77 (13)	C1—C6—C7—C8	-0.24 (13)
C9—N1—C1—C6	178.86 (12)	C5—C6—C7—C11	1.3 (2)
N1—C1—C2—C3	178.96 (11)	C1—C6—C7—C11	-177.94 (13)

C6—C1—C2—C3	-0.04 (18)	C1—N1—C8—C7	-0.95 (14)
C1—C2—C3—C4	0.56 (19)	C9—N1—C8—C7	-179.00 (12)
C2—C3—C4—C5	-0.6 (2)	C6—C7—C8—N1	0.73 (14)
C3—C4—C5—C6	0.0 (2)	C11—C7—C8—N1	178.60 (12)
C4—C5—C6—C1	0.47 (17)	C8—N1—C9—C10	11.7 (3)
C4—C5—C6—C7	-178.65 (12)	C1—N1—C9—C10	-166.00 (16)
C2—C1—C6—C5	-0.47 (17)	C8—C7—C11—O1	-178.09 (15)
N1—C1—C6—C5	-179.66 (10)	C6—C7—C11—O1	-0.7 (3)
C2—C1—C6—C7	178.87 (10)		

Hydrogen-bond geometry (Å, °)

<i>D</i> —H... <i>A</i>	<i>D</i> —H	H... <i>A</i>	<i>D</i> ... <i>A</i>	<i>D</i> —H... <i>A</i>
C9—H9...O1 ⁱ	0.93	2.51	3.390 (2)	159

Symmetry code: (i) $x-1/2, -y+1/2, z+1/2$.