

Acta Crystallographica Section E

Structure Reports

Online

ISSN 1600-5368

 Methyl 3-(1*H*-indol-3-yl)propanoate

Rui-Bin Hou and Dong-Feng Li*

 School of Chemistry and Life Science, Changchun University of Technology,
 Changchun 130012, People's Republic of China

Correspondence e-mail: lidongfeng@mail.ccut.edu.cn

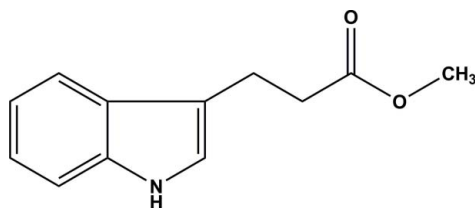
Received 13 July 2011; accepted 16 July 2011

 Key indicators: single-crystal X-ray study; $T = 296$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å;
 R factor = 0.052; wR factor = 0.138; data-to-parameter ratio = 17.5.

The molecule of the title compound, $\text{C}_{12}\text{H}_{13}\text{NO}_2$, adopts an essentially planar conformation (r.m.s. deviation = 0.057 Å). In the crystal, the molecules are linked by intermolecular $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds, generating chains along [201].

Related literature

For the biological activity of indole derivatives, see: Zeynep *et al.* (2005); Seefeld *et al.* (2003). For details of the synthesis, see: Pedras & Soledade (2006).



Experimental

Crystal data

$\text{C}_{12}\text{H}_{13}\text{NO}_2$
 $M_r = 203.23$
 Monoclinic, $P2_1/c$
 $a = 6.893$ (5) Å
 $b = 9.146$ (8) Å
 $c = 18.052$ (10) Å
 $\beta = 111.27$ (3)°

$V = 1060.5$ (13) Å³
 $Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 0.09$ mm⁻¹
 $T = 296$ K
 $0.46 \times 0.19 \times 0.18$ mm

Data collection

Rigaku R-Axis RAPID
 diffractometer
 Absorption correction: multi-scan
 (ABSCOR; Higashi, 1995)
 $T_{\min} = 0.961$, $T_{\max} = 0.984$

10015 measured reflections
 2414 independent reflections
 1509 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.048$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.052$
 $wR(F^2) = 0.138$
 $S = 1.04$
 2414 reflections
 138 parameters

1 restraint
 H-atom parameters constrained
 $\Delta\rho_{\max} = 0.16$ e Å⁻³
 $\Delta\rho_{\min} = -0.16$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
$\text{N1}-\text{H1}\cdots\text{O1}^{\dagger}$	0.95	2.08	2.972 (3)	157

 Symmetry code: (i) $x + 1, -y + \frac{3}{2}, z + \frac{1}{2}$

Data collection: *RAPID-AUTO* (Rigaku Corporation, 1998); cell refinement: *RAPID-AUTO*; data reduction: *CrystalStructure* (MSC & Rigaku, 2002); program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008) and *DIAMOND* (Brandenburg & Berndt, 2001); software used to prepare material for publication: *SHELXL97*.

The authors acknowledge financial support from the National Natural Science Foundation of Jilin Province (grant No. 20101548).

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: GK2396).

References

- Brandenburg, K. & Berndt, M. (2001). *DIAMOND*. Crystal Impact, Bonn, Germany.
- Higashi, T. (1995). *ABSCOR*. Rigaku Corporation, Tokyo, Japan.
- MSC & Rigaku (2002). *CrystalStructure*. MSC, The Woodlands, Texas, USA, and Rigaku Corporation, Tokyo, Japan.
- Pedras, M. & Soledade, C. (2006). *Med. Chem.* **14**, 4958–4979.
- Rigaku Corporation (1998). *RAPID-AUTO*. Rigaku Corporation, Tokyo, Japan.
- Seefeld, M. A., Miler, W. H., Newlander, K. A., Burgess, W. J., DeWolf, W. E. Jr, Elkins, P. A., Head, M. S., Jakas, D. R., Janson, C. A., Keller, P. M., Manley, P. J., Moore, T. D., Payne, D. J., Pearson, S., Polizzi, B. J., Qiu, X., Rittenhouse, S. F., Uzinskas, I. N., Wallis, N. G. & Huffman, W. F. (2003). *J. Med. Chem.* **46**, 1627–1635.
- Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.
- Zeynep, A. A., Tulay, C. & Sibel, S. (2005). *Med. Chem. Res.* **14**, 169–179.

supporting information

Acta Cryst. (2011). E67, o2121 [doi:10.1107/S1600536811028595]

Methyl 3-(1*H*-indol-3-yl)propanoate

Rui-Bin Hou and Dong-Feng Li

S1. Comment

Indole derivatives constitute an important class of therapeutic agents in medicinal chemistry including anticancer, antioxidant, antirheumatoidal and anti-HIV (Zeynep *et al.*, 2005; Seefeld *et al.*, 2003). We have recently synthesized some indole derivatives as histone deacetylase (HDAC) inhibitors with the precursor. In this paper, we report the crystal structure of the title compound.

The molecular structure of title compound, C₁₂H₁₃O₂N, as shown in Fig. 1, all bond lengths and angles are in the normal ranges. All non-hydrogen atoms except for O1 are nearly coplanar. In the crystal, the intermolecular N—H···O hydrogen bonds link the molecules into chains along the [201] direction.

S2. Experimental

The title compound was prepared according to the literature (Pedras & Soledade, 2006). Single crystals suitable for X-ray diffraction were prepared by slow evaporation method from a solution in dichloromethane/ petroleum (60–90 °C) at room temperature.

S3. Refinement

Carbon-bound H-atoms were placed in calculated positions (C—H 0.93 and 0.97 Å) and were included in the refinement in the riding model with $U_{\text{iso}}(\text{H}) = 1.5$ or $1.2 U_{\text{eq}}(\text{C})$. The N-bound H atom was located from a difference map and refined with the distance restraint N—H = 0.90 Å and $U_{\text{iso}}(\text{H}) = 1.5 U_{\text{eq}}(\text{N})$.

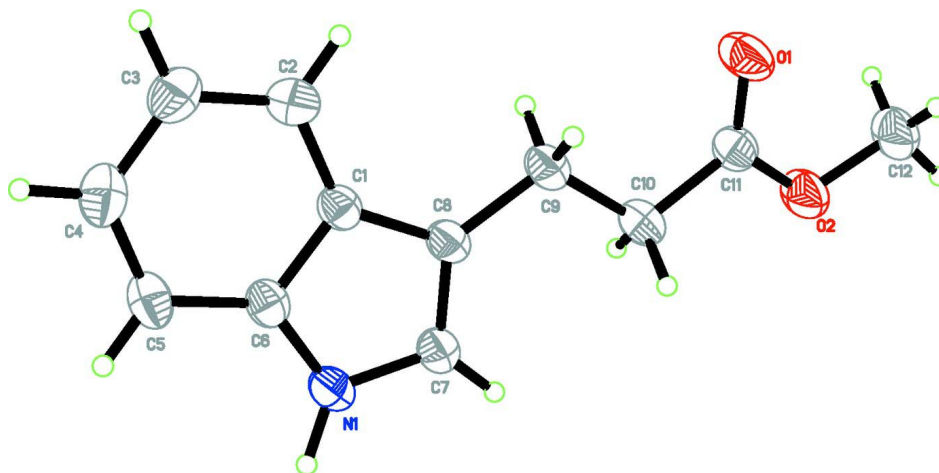
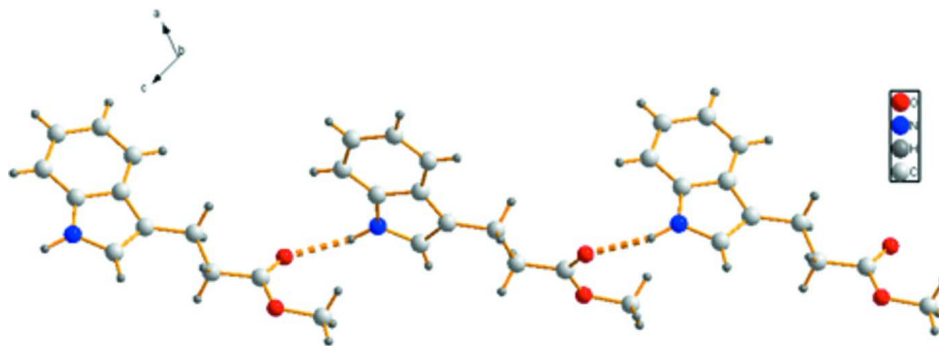


Figure 1

Molecular structure of the title compound with displacement ellipsoids drawn at the 30% probability level.

**Figure 2**

Hydrogen-bonded chain in the title compound. Dashed lines indicate hydrogen bonds.

Methyl 3-(1*H*-indol-3-yl)propanoate

Crystal data

$C_{12}H_{13}NO_2$

$M_r = 203.23$

Monoclinic, $P2_1/c$

Hall symbol: -P 2ybc

$a = 6.893$ (5) Å

$b = 9.146$ (8) Å

$c = 18.052$ (10) Å

$\beta = 111.27$ (3)°

$V = 1060.5$ (13) Å³

$Z = 4$

$F(000) = 432$

$D_x = 1.273$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 5849 reflections

$\theta = 3.2$ – 27.4 °

$\mu = 0.09$ mm⁻¹

$T = 296$ K

Block, colorless

$0.46 \times 0.19 \times 0.18$ mm

Data collection

Rigaku R-AXIS RAPID
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

ω scans

Absorption correction: multi-scan
(*ABSCOR*; Higashi, 1995)

$T_{\min} = 0.961$, $T_{\max} = 0.984$

10015 measured reflections

2414 independent reflections

1509 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.048$

$\theta_{\max} = 27.5$ °, $\theta_{\min} = 3.2$ °

$h = -8 \rightarrow 8$

$k = -11 \rightarrow 11$

$l = -23 \rightarrow 23$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.052$

$wR(F^2) = 0.138$

$S = 1.04$

2414 reflections

138 parameters

1 restraint

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

$w = 1/[\sigma^2(F_o^2) + (0.0614P)^2 + 0.1289P]$

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

$(\Delta/\sigma)_{\max} < 0.001$

$\Delta\rho_{\max} = 0.16$ e Å⁻³

$\Delta\rho_{\min} = -0.16$ e Å⁻³

Extinction correction: *SHELXL*,

$F_c^* = kF_c[1 + 0.001x F_c^2 \lambda^3 / \sin(2\theta)]^{-1/4}$

Extinction coefficient: 0.036 (5)

Special details

Experimental. (See detailed section in the paper)

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 (\AA^2)

	x	y	z	$U_{\text{iso}}^*/U_{\text{eq}}$
O1	-0.1892 (2)	0.69967 (17)	0.43997 (8)	0.0762 (5)
O2	-0.3213 (2)	0.88674 (17)	0.48330 (8)	0.0760 (5)
N1	0.5979 (2)	0.80147 (17)	0.76433 (8)	0.0566 (4)
H1	0.6448	0.8259	0.8192	0.085*
C1	0.5726 (3)	0.65764 (19)	0.66104 (9)	0.0469 (4)
C2	0.6478 (3)	0.5573 (2)	0.61952 (11)	0.0585 (5)
H2	0.5628	0.5253	0.5693	0.070*
C3	0.8477 (3)	0.5065 (2)	0.65345 (13)	0.0668 (6)
H3	0.8984	0.4403	0.6258	0.080*
C4	0.9761 (3)	0.5526 (2)	0.72877 (13)	0.0671 (6)
H4	1.1109	0.5161	0.7507	0.080*
C5	0.9079 (3)	0.6506 (2)	0.77134 (11)	0.0598 (5)
H5	0.9941	0.6810	0.8217	0.072*
C6	0.7059 (3)	0.70276 (19)	0.73679 (10)	0.0481 (4)
C7	0.4000 (3)	0.8150 (2)	0.70877 (10)	0.0546 (5)
H7	0.2961	0.8736	0.7145	0.066*
C8	0.3779 (3)	0.7306 (2)	0.64415 (9)	0.0491 (5)
C9	0.1912 (3)	0.7142 (2)	0.56905 (10)	0.0606 (5)
H9A	0.1467	0.6129	0.5639	0.073*
H9B	0.2316	0.7372	0.5242	0.073*
C10	0.0095 (3)	0.8096 (2)	0.56507 (10)	0.0558 (5)
H10A	-0.0319	0.7868	0.6097	0.067*
H10B	0.0529	0.9111	0.5699	0.067*
C11	-0.1734 (3)	0.7903 (2)	0.48976 (10)	0.0532 (5)
C12	-0.5084 (3)	0.8803 (3)	0.41298 (12)	0.0820 (7)
H12A	-0.4753	0.9036	0.3671	0.123*
H12B	-0.6080	0.9494	0.4178	0.123*
H12C	-0.5661	0.7836	0.4074	0.123*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
O1	0.0724 (10)	0.0813 (11)	0.0522 (7)	0.0196 (8)	-0.0047 (7)	-0.0124 (7)
O2	0.0590 (9)	0.0844 (11)	0.0673 (8)	0.0192 (8)	0.0020 (7)	-0.0161 (7)
N1	0.0537 (9)	0.0640 (10)	0.0445 (8)	-0.0005 (8)	0.0087 (7)	-0.0074 (7)

C1	0.0477 (10)	0.0480 (10)	0.0424 (8)	-0.0038 (8)	0.0132 (8)	0.0026 (7)
C2	0.0677 (13)	0.0552 (12)	0.0516 (10)	-0.0013 (10)	0.0204 (9)	-0.0030 (8)
C3	0.0675 (14)	0.0577 (13)	0.0807 (14)	0.0070 (10)	0.0338 (11)	-0.0018 (10)
C4	0.0520 (12)	0.0571 (13)	0.0878 (14)	0.0035 (10)	0.0202 (11)	0.0053 (11)
C5	0.0475 (11)	0.0566 (12)	0.0635 (11)	-0.0053 (9)	0.0060 (9)	0.0001 (9)
C6	0.0458 (10)	0.0459 (10)	0.0486 (9)	-0.0054 (8)	0.0124 (8)	0.0030 (8)
C7	0.0476 (11)	0.0620 (12)	0.0485 (9)	0.0046 (9)	0.0107 (8)	-0.0029 (8)
C8	0.0487 (10)	0.0542 (11)	0.0401 (9)	-0.0019 (8)	0.0109 (8)	-0.0001 (7)
C9	0.0529 (11)	0.0744 (14)	0.0440 (9)	0.0054 (10)	0.0050 (8)	-0.0041 (9)
C10	0.0574 (12)	0.0549 (11)	0.0464 (9)	-0.0020 (9)	0.0084 (8)	0.0004 (8)
C11	0.0542 (11)	0.0548 (11)	0.0460 (9)	0.0027 (9)	0.0126 (8)	0.0045 (8)
C12	0.0555 (13)	0.0987 (19)	0.0722 (13)	0.0191 (12)	-0.0002 (11)	-0.0103 (12)

Geometric parameters (Å, °)

O1—C11	1.198 (2)	C5—C6	1.388 (3)
O2—C11	1.321 (2)	C5—H5	0.9300
O2—C12	1.446 (2)	C7—C8	1.361 (3)
N1—C6	1.373 (2)	C7—H7	0.9300
N1—C7	1.375 (2)	C8—C9	1.500 (2)
N1—H1	0.9498	C9—C10	1.507 (3)
C1—C2	1.398 (3)	C9—H9A	0.9700
C1—C6	1.404 (2)	C9—H9B	0.9700
C1—C8	1.429 (3)	C10—C11	1.491 (2)
C2—C3	1.371 (3)	C10—H10A	0.9700
C2—H2	0.9300	C10—H10B	0.9700
C3—C4	1.392 (3)	C12—H12A	0.9600
C3—H3	0.9300	C12—H12B	0.9600
C4—C5	1.370 (3)	C12—H12C	0.9600
C4—H4	0.9300		
C11—O2—C12	117.67 (16)	C7—C8—C1	106.12 (15)
C6—N1—C7	108.66 (15)	C7—C8—C9	128.62 (17)
C6—N1—H1	120.5	C1—C8—C9	125.26 (16)
C7—N1—H1	127.7	C8—C9—C10	114.27 (16)
C2—C1—C6	118.42 (17)	C8—C9—H9A	108.7
C2—C1—C8	133.97 (16)	C10—C9—H9A	108.7
C6—C1—C8	107.61 (16)	C8—C9—H9B	108.7
C3—C2—C1	119.42 (18)	C10—C9—H9B	108.7
C3—C2—H2	120.3	H9A—C9—H9B	107.6
C1—C2—H2	120.3	C11—C10—C9	112.84 (16)
C2—C3—C4	120.9 (2)	C11—C10—H10A	109.0
C2—C3—H3	119.5	C9—C10—H10A	109.0
C4—C3—H3	119.5	C11—C10—H10B	109.0
C5—C4—C3	121.4 (2)	C9—C10—H10B	109.0
C5—C4—H4	119.3	H10A—C10—H10B	107.8
C3—C4—H4	119.3	O1—C11—O2	122.57 (17)
C4—C5—C6	117.73 (18)	O1—C11—C10	125.61 (18)

C4—C5—H5	121.1	O2—C11—C10	111.82 (16)
C6—C5—H5	121.1	O2—C12—H12A	109.5
N1—C6—C5	130.60 (17)	O2—C12—H12B	109.5
N1—C6—C1	107.26 (15)	H12A—C12—H12B	109.5
C5—C6—C1	122.14 (17)	O2—C12—H12C	109.5
C8—C7—N1	110.32 (16)	H12A—C12—H12C	109.5
C8—C7—H7	124.8	H12B—C12—H12C	109.5
N1—C7—H7	124.8		
C6—C1—C2—C3	-0.2 (3)	N1—C7—C8—C1	1.2 (2)
C8—C1—C2—C3	-179.77 (19)	N1—C7—C8—C9	-178.59 (18)
C1—C2—C3—C4	0.5 (3)	C2—C1—C8—C7	179.6 (2)
C2—C3—C4—C5	-0.4 (3)	C6—C1—C8—C7	-0.1 (2)
C3—C4—C5—C6	-0.1 (3)	C2—C1—C8—C9	-0.6 (3)
C7—N1—C6—C5	-178.76 (18)	C6—C1—C8—C9	179.73 (17)
C7—N1—C6—C1	1.8 (2)	C7—C8—C9—C10	2.6 (3)
C4—C5—C6—N1	-179.00 (18)	C1—C8—C9—C10	-177.13 (17)
C4—C5—C6—C1	0.4 (3)	C8—C9—C10—C11	-179.92 (16)
C2—C1—C6—N1	179.23 (15)	C12—O2—C11—O1	0.1 (3)
C8—C1—C6—N1	-1.06 (19)	C12—O2—C11—C10	-179.83 (17)
C2—C1—C6—C5	-0.3 (3)	C9—C10—C11—O1	7.5 (3)
C8—C1—C6—C5	179.44 (16)	C9—C10—C11—O2	-172.55 (16)
C6—N1—C7—C8	-1.9 (2)		

Hydrogen-bond geometry (Å, °)

<i>D</i> —H \cdots <i>A</i>	<i>D</i> —H	H \cdots <i>A</i>	<i>D</i> \cdots <i>A</i>	<i>D</i> —H \cdots <i>A</i>
N1—H1 \cdots O1 ⁱ	0.95	2.08	2.972 (3)	157

Symmetry code: (i) $x+1, -y+3/2, z+1/2$.