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The effect of hydrogen bonding on the conformations of 2-(1*H*-indol-3-yl)-2-oxoacetamide and 2-(1*H*-indol-3-yl)-*N*,*N*-dimethyl-2-oxoacetamide

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In the title compounds, $C_{10}H_8N_2O_2$, (I), and $C_{12}H_{12}N_2O_2$, (II), the two carbonyl groups are oriented with torsion angles of -149.3 (3) and -88.55 (15)°, respectively. The single-bond distances linking the two carbonyl groups are 1.528 (4) and 1.5298 (17) Å, respectively. In (I), the molecules are linked by an elaborate system of N-H···O hydrogen bonds, which form adjacent $R_2^2(8)$ and $R_2^4(8)$ ring motifs to generate a ladder-like construct. Adjacent ladders are further linked by N-H···O hydrogen bonds to build a three-dimensional network. The hydrogen bonding in (II) is far simpler, consisting of helical chains of N-H···O-linked molecules that follow the 2_1 screw of the *b* axis. It is the presence of an elaborate hydrogen-bonding system in the crystal structure of (I) that leads to the different torsion angle for the orientation of the two adjacent carbonyl groups from that in (II).

Comment

The title compounds, 2-(1H-indol-3-yl)-2-oxoacetamide, (I), and 2-(1H-indol-3-yl)-N,N-dimethyl-2-oxoacetamide, (II), are synthetic intermediates in the preparation of tryptamine and N,N-dimethyltryptamine, and are prepared by treating indole with oxalyl chloride followed by quenching with aqueous ammonia or dimethylamine, respectively. The products were characterized by spectroscopic analysis and their X-ray crystal structure determinations were carried out to ascertain the conformations of these molecules, especially the orientations of the carbonyl groups, as 1,2-diones have been shown to be potent inhibitors of carboxylesterases (Hyatt et al., 2007). The inhibitory properties of ethane-1,2-diones toward carboxylesterases are such that, when the carbonyl O atoms are ciscoplanar, the compounds demonstrate specificity for human liver carboxylesterase. Conversely, when the dione O atoms are not planar (or are *trans*-coplanar), the compounds are more potent at human intestinal carboxylesterase inhibition (Hyatt *et al.*, 2007). Thus, the title compounds and their analogues were screened for their inhibition of human carboxylesterases and it was planned to determine their conformations to compare with the observation reported for ethane-1,2-diones.



X-ray crystallography confirmed the molecular structures and atom connectivities of (I) and (II) (Figs. 1 and 3), and selected geometric parameters are listed in Tables 1 and 3. The two carbonyl groups are in a *trans* orientation, with torsion angles O1-C9-C10-O2 = -149.3 (3)° in (I) and -88.55 (15)° in (II). The deviation from 180° may be due to repulsive interactions between nonbonded electron pairs on atoms O1 and O2. However, in (I) the extensive hydrogen bonding introduces rigidity into the molecule. Since in both compounds the C9=O1 groups are coplanar with the indole nucleus, there is extended conjugation from atom O1 through to the indole ring. This is evident from the shortening of the



Figure 1

A view of the molecule of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level.





A packing diagram for (I), viewed down the c axis. Dashed lines indicate hydrogen bonds. H atoms not involved in these interactions have been omitted for clarity.



Figure 3

A view of the molecule of (II), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level.





A packing diagram for (II), viewed down the b axis. Dashed lines indicate hydrogen bonds. H atoms not involved in these interactions have been omitted for clarity.

C8–C9 bonds [1.421 (4) Å in (I) and 1.4380 (17) Å in (II)] bonds compared with the standard value for a single bond connecting a C_{ar} atom to a Csp^2 atom [1.470 (15) Å; Allen *et al.*, 1987]. Possibly due to repulsive interactions between nonbonded electron pairs, the C9–C10 bonds are stretched, resulting in bond lengths of 1.528 (4) and 1.5298 (17) Å in (I) and (II), respectively. These are longer than expected, the characteristic value for a Csp^2-Csp^2 bond being 1.50 Å (Zukerman-Schpector *et al.*, 1994). The C10–N2 bond lengths [1.323 (4) Å in (I) and 1.3287 (16) Å in (II)], and the bond angles around the N2 atoms, suggest that the lone pairs of electrons on N2 undergo delocalization, bestowing doublebond character on the C10–N2 bonds.

The packing of compounds (I) and (II), as viewed down the c and b axes, are illustrated in Figs. 2 and 4, respectively.

Amides undergo extensive hydrogen bonding, but in (I), in addition to primary amide functionality, there is a hydrogenbond-accepting carbonyl group and a variable glyoxylamide torsion angle. In (I), molecules are linked by an elaborate system of $N-H\cdots O$ hydrogen bonds (entries 1 and 2 in Table 2) form adjacent $R_2^2(8)$ and $R_2^4(8)$ ring motifs (Bernstein *et al.*, 1995) to generate a ladder-like construct. Adjacent ladders are further linked by additional $N-H\cdots O$ hydrogen bonds (entry 3 in Table 2) to build a three-dimensional network. The hydrogen bonding in (II) is far simpler, consisting of helical chains of $N-H\cdots O$ -linked molecules that follow the 2_1 screw of the *b* axis (Table 4).

Experimental

The title compounds were prepared according to a previously reported procedure (Da Settimo *et al.*, 1996). Recrystallization from ethyl acetate afforded (I) and (II) as pale-yellow and colourless crystalline products which were suitable for X-ray analysis. Spectroscopic data for (I), ¹H NMR (300 MHz, DMSO-*d*₆): δ 7.22–7.25 (*m*, 2H), 7.49–7.52 (*m*, 1H), 7.70 (*sb*, 1H), 8.06 (*sb*, 1H), 8.19–8.22 (*m*, 1H), 8.67 (*d*, 1H), 12.17 (*sb*, 1H); ¹³C NMR (75 MHz, DMSO-*d*₆): δ 111.95, 112.41, 121.10, 122.32, 123.22, 125.98, 136.11, 138.10, 165.80, 182.72. Spectroscopic data for (II), ¹H NMR (300 MHz, DMSO-*d*₆): δ 2.93 (*s*, 3H), 3.01 (*s*, 3H), 7.23–7.32 (*m*, 2H), 7.51–7.57 (*m*, 1H), 8.10 (*dd*, 1H), 8.14 (*s*, 1H), 12.17 (*sb*, 1H); ¹³C NMR (75 MHz, DMSO-*d*₆): δ 33.41, 36.81, 112.52, 112.83, 120.71, 122.38, 123.38, 124.75, 136.73, 136.83, 167.11, 186.44.

Compound (I)

Crystal data

 $C_{10}H_8N_2O_2$ $M_r = 188.18$ Monoclinic, $P2_1/c$ a = 5.1253 (4) Å b = 19.4363 (14) Å c = 8.4953 (6) Å $\beta = 93.251$ (3)°

Data collection

Bruker X8 Proteum diffractometer Absorption correction: multi-scan [SADABS (Bruker, 2006) and XABS2 (Parkin et al., 1995)] T_{min} = 0.744, T_{max} = 0.949

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.058$ $wR(F^2) = 0.158$ S = 1.241518 reflections

Compound (II)

Crystal data $C_{12}H_{12}N_2O_2$ $M_r = 216.24$ Monoclinic, $P2_1/c$ a = 10.1813 (2) Å b = 6.0048 (1) Å c = 18.0625 (4) Å $\beta = 97.7222$ (8)° $V = 844.91 (11) Å^{3}$ Z = 4 Cu K\alpha radiation $\mu = 0.88 \text{ mm}^{-1}$ T = 90 K 0.30 × 0.08 × 0.06 mm

10887 measured reflections 1518 independent reflections 1481 reflections with $I > 2\sigma(I)$ $R_{\text{int}} = 0.042$

128 parameters H-atom parameters constrained
$$\begin{split} &\Delta\rho_{max}=0.33\ e\ \text{\AA}^{-3}\\ &\Delta\rho_{min}=-0.33\ e\ \text{\AA}^{-3} \end{split}$$

 $V = 1094.27 (4) Å^{3}$ Z = 4 Mo K\alpha radiation $\mu = 0.09 \text{ mm}^{-1}$ T = 90 K 0.65 × 0.25 × 0.15 mm

Table 1Selected geometric parameters (Å, $^{\circ}$) for (I).

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N2-C10	1.323 (4)	C8-C9	1.421 (4)
O2-C10	1.232 (3)	C9-C10	1.528 (4)
C1-C8-C9	126.9 (3)	O2-C10-N2	124.1 (3)
01-C9-C8	124.1 (3)	O2-C10-C9	122.3 (2)
O1-C9-C10	117.1 (3)	N2-C10-C9	113.5 (2)
C1-C8-C9-O1	-178.2 (3)	O1-C9-C10-O2	-149.3 (3)

Table 2

Hydrogen-bond geometry (Å, $^{\circ}$) for (I).

$D - H \cdot \cdot \cdot A$	$D-{\rm H}$	$H \cdot \cdot \cdot A$	$D \cdots A$	$D - H \cdots A$
$N1-H1\cdotsO1^{i}$ $N2-H2A\cdotsO2^{ii}$ $N2-H2B\cdotsO2^{iii}$	0.88 0.88 0.88	1.92 2.08 2.14	2.768 (3) 2.926 (3) 2.898 (3)	160 161 144

Symmetry codes: (i) x - 1, $-y + \frac{3}{2}$, $z + \frac{1}{2}$; (ii) -x + 1, -y + 1, -z + 1; (iii) x + 1, y, z.

Data collection

Nonius KappaCCD area-detector	4780 measured reflections
diffractometer	2515 independent reflections
Absorption correction: multi-scan	2031 reflections with $I > 2\sigma(I)$
(SCALEPACK; Otwinowski &	$R_{\rm int} = 0.033$
Minor, 1997)	
$T_{\rm min} = 0.943, T_{\rm max} = 0.986$	

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.040$	147 parameters
$wR(F^2) = 0.112$	H-atom parameters constrained
S = 1.06	$\Delta \rho_{\rm max} = 0.23 \ {\rm e} \ {\rm \AA}^{-3}$
2515 reflections	$\Delta \rho_{\rm min} = -0.20 \ {\rm e} \ {\rm \AA}^{-3}$

H atoms were found in difference Fourier maps and subsequently placed in idealized positions, with C-H = 0.95 (aromatic), 0.98 (methyl) or 0.88 Å (NH₂), and with $U_{iso}(H) = 1.5U_{eq}(C)$ for methyl H atoms or $1.2U_{eq}(C,N)$ otherwise.

Data collection: *APEX2* (Bruker, 2006) for (I); *COLLECT* (Nonius, 1999) for (II). Cell refinement: *APEX2* for (I); *SCALE-PACK* (Otwinowski & Minor, 1997) for (II). Data reduction: *APEX2* for (I); *DENZO-SMN* (Otwinowski & Minor, 1997) for (II). For both compounds, program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *XP* in *SHELXTL* (Sheldrick, 2008);

Table 3					
Selected	geometric parameters (Å,	°)	for	(II)	۱.

N2-C10	1.3287 (16)	C8-C9	1.4380 (17)
O2-C10	1.2431 (15)	C9-C10	1.5298 (17)
C1-C8-C9	125.50 (11)	O2-C10-N2	124.03 (11)
01-C9-C8	124.97 (12)	O2-C10-C9	118.05 (11)
O1-C9-C10	118.72 (11)	N2-C10-C9	117.91 (10)
C1-C8-C9-O1	175.49 (12)	O1-C9-C10-O2	-88.55 (15)

Table 4

Hydrogen-bond geometry (Å, °) for (II).

$D - H \cdot \cdot \cdot A$	D-H	$H \cdots A$	$D \cdot \cdot \cdot A$	$D - \mathbf{H} \cdot \cdot \cdot A$
$N1 - H1A \cdots O2^{i}$	0.88	1.92	2.7885 (14)	170
Symmetry code: (i) -	$-x + 1, y - \frac{1}{2}, -$	$z + \frac{1}{2}$.		

2008); software used to prepare material for publication: *SHELXL97* and local procedures.

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Supplementary data for this paper are available from the IUCr electronic archives (Reference: YP3017). Services for accessing these data are described at the back of the journal.

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supplementary materials

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(I) 2-(1*H*-indol-3-yl)-2-oxoacetamide

Crystal data

C₁₀H₈N₂O₂ $M_r = 188.18$ Monoclinic, $P2_1/c$ Hall symbol: -P 2ybc a = 5.1253 (4) Å b = 19.4363 (14) Å c = 8.4953 (6) Å $\beta = 93.251$ (3)° V = 844.91 (11) Å³ Z = 4

Data collection

Bruker X8 Proteum diffractometer Radiation source: fine-focus rotating anode Graded multilayer optics monochromator Detector resolution: 18 pixels mm⁻¹ φ and ω scans Absorption correction: multi-scan [*SADABS* (Bruker, 2006) and *XABS2* (Parkin *et al.*, 1995)]

Refinement

Refinement on F^2 Least-squares matrix: full $R[F^2 > 2\sigma(F^2)] = 0.058$ $wR(F^2) = 0.158$ S = 1.241518 reflections 128 parameters 0 restraints Primary atom site location: structure-invariant direct methods Secondary atom site location: difference Fourier map F(000) = 392 $D_x = 1.479 \text{ Mg m}^{-3}$ Cu K\alpha radiation, $\lambda = 1.54178 \text{ Å}$ Cell parameters from 8721 reflections $\theta = 4.6-69.0^{\circ}$ $\mu = 0.88 \text{ mm}^{-1}$ T = 90 KNeedle, colourless $0.30 \times 0.08 \times 0.06 \text{ mm}$

 $T_{\min} = 0.744, T_{\max} = 0.949$ 10887 measured reflections 1518 independent reflections 1481 reflections with $I > 2\sigma(I)$ $R_{int} = 0.042$ $\theta_{\max} = 68.9^{\circ}, \theta_{\min} = 4.6^{\circ}$ $h = -4 \rightarrow 6$ $k = -23 \rightarrow 23$ $l = -10 \rightarrow 10$

Hydrogen site location: inferred from neighbouring sites H-atom parameters constrained $w = 1/[\sigma^2(F_o^2) + (0.037P)^2 + 1.8838P]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{max} < 0.001$ $\Delta\rho_{max} = 0.33$ e Å⁻³ $\Delta\rho_{min} = -0.33$ e Å⁻³ Extinction correction: *SHELXL97* (Sheldrick, 2008), Fc*=kFc[1+0.001xFc^2\lambda^3/sin(2\theta)]^{-1/4} Extinction coefficient: 0.0033 (9)

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 $F^2 > 2\sigma(F^2)$ 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.

	X	у	Ζ	$U_{\rm iso}$ */ $U_{\rm eq}$
N1	0.0782 (5)	0.77152 (13)	0.5558 (3)	0.0227 (6)
H1	-0.0529	0.7766	0.6173	0.027*
N2	0.7272 (5)	0.56011 (12)	0.4103 (3)	0.0219 (6)
H2A	0.7242	0.5163	0.4362	0.026*
H2B	0.8761	0.5800	0.3904	0.026*
01	0.7253 (4)	0.68365 (10)	0.2684 (2)	0.0221 (5)
O2	0.2916 (4)	0.57240 (10)	0.4272 (3)	0.0231 (5)
C1	0.1826 (5)	0.71168 (15)	0.5181 (3)	0.0195 (6)
H1A	0.1282	0.6681	0.5546	0.023*
C2	0.2054 (6)	0.82452 (15)	0.4845 (3)	0.0207 (7)
C3	0.1592 (6)	0.89462 (16)	0.4918 (4)	0.0274 (7)
H3	0.0254	0.9129	0.5524	0.033*
C4	0.3164 (6)	0.93666 (16)	0.4069 (4)	0.0297 (8)
H4	0.2880	0.9850	0.4068	0.036*
C5	0.5163 (6)	0.90938 (16)	0.3211 (4)	0.0270 (7)
Н5	0.6249	0.9396	0.2663	0.032*
C6	0.5597 (6)	0.83963 (15)	0.3143 (4)	0.0225 (7)
H6	0.6958	0.8218	0.2548	0.027*
C7	0.4006 (5)	0.79562 (15)	0.3960 (3)	0.0186 (6)
C8	0.3834 (5)	0.72223 (15)	0.4173 (3)	0.0183 (6)
С9	0.5462 (5)	0.67120 (15)	0.3542 (3)	0.0185 (6)
C10	0.5078 (5)	0.59616 (15)	0.4013 (3)	0.0192 (6)

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters $(Å^2)$

Atomic displacement parameters $(Å^2)$

	I 711	I 122	I 733	I /12	<i>T</i> /13	I 723
	U	U	U	U	0	U
N1	0.0145 (12)	0.0325 (14)	0.0213 (13)	0.0038 (10)	0.0036 (10)	-0.0025 (10)
N2	0.0094 (11)	0.0211 (13)	0.0356 (15)	-0.0008 (9)	0.0044 (10)	0.0028 (10)
O1	0.0138 (10)	0.0286 (11)	0.0247 (11)	-0.0003 (8)	0.0072 (8)	0.0019 (8)
02	0.0120 (10)	0.0239 (11)	0.0335 (12)	-0.0002 (8)	0.0037 (8)	0.0031 (9)
C1	0.0142 (14)	0.0269 (15)	0.0175 (14)	0.0009 (11)	0.0007 (11)	0.0001 (11)
C2	0.0155 (14)	0.0276 (16)	0.0186 (15)	0.0018 (11)	-0.0028 (11)	-0.0035 (11)
C3	0.0221 (16)	0.0295 (17)	0.0301 (18)	0.0043 (13)	-0.0037 (13)	-0.0079 (13)
C4	0.0274 (17)	0.0231 (16)	0.0375 (19)	0.0039 (13)	-0.0071 (14)	-0.0030 (13)
C5	0.0239 (16)	0.0261 (16)	0.0302 (18)	-0.0039 (12)	-0.0049 (13)	0.0023 (13)
C6	0.0167 (15)	0.0268 (16)	0.0237 (16)	-0.0012 (12)	-0.0016 (12)	0.0005 (12)
C7	0.0130 (13)	0.0253 (15)	0.0171 (14)	0.0009 (11)	-0.0024 (11)	-0.0001 (11)

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C8	0.0124 (13)	0.0229 (14)	0.0195 (14)	0.0015 (11)	0.0011 (11)	-0.0003 (11)
C9	0.0118 (13)	0.0238 (15)	0.0200 (15)	-0.0011 (11)	0.0009 (11)	0.0007 (11)
C10	0.0128 (14)	0.0245 (15)	0.0206 (15)	-0.0003 (11)	0.0028 (11)	-0.0010 (11)
Geomet	ric parameters (Å	, <i>o</i>)				
N1—C1		1.327 ((4)	C3—C4		1.379 (5)
N1-C2	2	1.378 ((4)	С3—Н3		0.9500
N1—H1	l	0.8800		C4—C5		1.395 (5)
N2—C1	0	1.323 ((4)	C4—H4		0.9500
N2—H2	2A	0.8800		C5—C6		1.376 (4)
N2—H2	2B	0.8800		С5—Н5		0.9500
01—C9)	1.228 ((3)	С6—С7		1.394 (4)
02—C1	0	1.232 ((3)	С6—Н6		0.9500
C1—C8	3	1.391 ((4)	С7—С8		1.441 (4)
С1—Н1	A	0.9500		С8—С9		1.421 (4)
C2—C3	3	1.385 ((4)	C9—C10		1.528 (4)
C2—C7	1	1.403 ((4)			
C1—N1	—C2	110.0 (2)	C6—C5—C4		121.5 (3)
C1—N1	—H1	125.0		С6—С5—Н5		119.3
C2—N1	—H1	125.0		С4—С5—Н5		119.3
C10—N	I2—H2A	120.0		C5—C6—C7		118.9 (3)
C10—N	12—H2B	120.0		С5—С6—Н6		120.6
H2A—I	N2—H2B	120.0		С7—С6—Н6		120.6
N1—C1	—C8	110.0 (3)	C6—C7—C2		118.5 (3)
N1—C1	—H1A	125.0		С6—С7—С8		135.4 (3)
C8-C1	—H1A	125.0		С2—С7—С8		106.2 (2)
N1-C2	2—С3	129.1 ((3)	C1—C8—C9		126.9 (3)
N1-C2	2—С7	107.8 ((3)	C1—C8—C7		106.0 (2)
C3—C2	2—С7	123.1 (3)	C9—C8—C7 127.0 (3)		127.0 (3)
C4—C3	G—C2	117.0 (3)	01—C9—C8 124.1 (3)		
C4—C3	9—Н3	121.5		O1—C9—C10	117.1 (3)	
C2—C3	9—Н3	121.5		C8—C9—C10		118.7 (2)
C3—C4	⊢C5	121.1 ((3)	O2—C10—N2		124.1 (3)
C3—C4	H4	119.5		O2—C10—C9		122.3 (2)
C5—C4	—H4	119.5		N2—C10—C9		113.5 (2)
C2—N1		-0.9 (3	5)	N1—C1—C8—C9)	178.3 (3)
C1—N1		179.8 ((3)	N1-C1-C8-C7	7	1.0 (3)
C1—N1		0.4 (3)		C6—C7—C8—C1		178.4 (3)
N1-C2	2—C3—C4	-179.5	(3)	C2—C7—C8—C1		-0.7 (3)
C7—C2	2—C3—C4	-0.2 (4)	C6—C7—C8—C9)	1.0 (5)
C2—C3	G-C4-C5	-1.5 (5	5)	C2—C7—C8—C9)	-178.1 (3)
C3—C4	└──C5──C6	1.9 (5)		C1—C8—C9—O1	l	-178.2 (3)
C4—C5	Б—С6—С7	-0.4 (5	5)	С7—С8—С9—О1	l	-1.3 (5)
С5—Сб	б—С7—С2	-1.2 (4	-)	C1—C8—C9—C1	0	-1.4 (4)
С5—Сб	б—С7—С8	179.8 ((3)	C7—C8—C9—C1	0	175.5 (3)
N1-C2	2—С7—С6	-179.0	(3)	O1—C9—C10—C	02	-149.3 (3)
C3—C2	2—С7—С6	1.5 (4)		C8—C9—C10—C	02	33.7 (4)

N1-C2-C7-C8	0.2 (3)	O1-C9-C10-N2	30.4 (4)
C3—C2—C7—C8	-179.2 (3)	C8—C9—C10—N2	-146.6 (3)

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
N1—H1···O1 ⁱ	0.88	1.92	2.768 (3)	160
N2—H2A····O2 ⁱⁱ	0.88	2.08	2.926 (3)	161
N2—H2 <i>B</i> ···O2 ⁱⁱⁱ	0.88	2.14	2.898 (3)	144

4780 measured reflections 2515 independent reflections 2031 reflections with $I > 2\sigma(I)$

 $\theta_{\rm max} = 27.5^{\circ}, \ \theta_{\rm min} = 2.0^{\circ}$

 $R_{\rm int} = 0.033$

 $h = -13 \rightarrow 13$ $k = -7 \rightarrow 7$ $l = -23 \rightarrow 23$

Symmetry codes: (i) *x*-1, -*y*+3/2, *z*+1/2; (ii) -*x*+1, -*y*+1, -*z*+1; (iii) *x*+1, *y*, *z*.

(II) 2-(1H-indol-3-yl)-N,N-dimethyl-2-oxoacetamide

Crystal data

$C_{12}H_{12}N_2O_2$	F(000) = 456
$M_r = 216.24$	$D_{\rm x} = 1.313 {\rm ~Mg} {\rm ~m}^{-3}$
Monoclinic, $P2_1/c$	Mo <i>K</i> α radiation, $\lambda = 0.71073$ Å
Hall symbol: -P 2ybc	Cell parameters from 2759 reflections
a = 10.1813 (2) Å	$\theta = 1.0-27.5^{\circ}$
b = 6.0048 (1) Å	$\mu = 0.09 \text{ mm}^{-1}$
c = 18.0625 (4) Å	T = 90 K
$\beta = 97.7222 \ (8)^{\circ}$	Rod, colourless
$V = 1094.27 (4) Å^3$	$0.65 \times 0.25 \times 0.15 \text{ mm}$
Z = 4	

Data collection

Nonius KappaCCD area-detector
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
Detector resolution: 18 pixels mm ⁻¹
ω scans at fixed $\chi = 55^{\circ}$
Absorption correction: multi-scan
SCALEPACK (Otwinowski & Minor, 1997)
$T_{\min} = 0.943, T_{\max} = 0.986$

Refinement

Secondary atom site location: difference Fourier
map
Hydrogen site location: inferred from
neighbouring sites
H-atom parameters constrained
$w = 1/[\sigma^2(F_o^2) + (0.0561P)^2 + 0.2642P]$
where $P = (F_o^2 + 2F_c^2)/3$
$(\Delta/\sigma)_{\rm max} < 0.001$
$\Delta \rho_{\rm max} = 0.23 \ {\rm e} \ {\rm \AA}^{-3}$
$\Delta \rho_{\min} = -0.20 \text{ e} \text{ Å}^{-3}$

Special details

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

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

	x	у	Ζ	$U_{\rm iso}$ */ $U_{\rm eq}$
C1	0.46067 (12)	0.5029 (2)	0.30948 (7)	0.0250 (3)
H1	0.5263	0.4004	0.2983	0.030*
N1	0.32928 (10)	0.47893 (19)	0.28893 (6)	0.0262 (3)
H1A	0.2914	0.3676	0.2625	0.031*
N2	0.77326 (10)	0.48231 (18)	0.42133 (6)	0.0253 (3)
01	0.63689 (9)	0.95880 (17)	0.41126 (6)	0.0341 (3)
O2	0.78438 (8)	0.65588 (16)	0.31061 (5)	0.0295 (2)
C2	0.26301 (12)	0.6570(2)	0.31595 (6)	0.0229 (3)
C3	0.12758 (12)	0.6955 (2)	0.31215 (7)	0.0268 (3)
H3	0.0643	0.5964	0.2866	0.032*
C4	0.08930 (13)	0.8841 (3)	0.34723 (7)	0.0310 (3)
H4	-0.0024	0.9150	0.3463	0.037*
C5	0.18277 (13)	1.0309 (2)	0.38423 (8)	0.0317 (3)
Н5	0.1530	1.1600	0.4074	0.038*
C6	0.31755 (12)	0.9925 (2)	0.38792 (7)	0.0264 (3)
H6	0.3802	1.0936	0.4130	0.032*
C7	0.35877 (11)	0.8014 (2)	0.35382 (6)	0.0217 (3)
C8	0.48596 (11)	0.6990 (2)	0.34928 (6)	0.0218 (3)
C9	0.61495 (11)	0.7791 (2)	0.37998 (7)	0.0229 (3)
C10	0.73240 (11)	0.6301 (2)	0.36848 (7)	0.0224 (3)
C11	0.70814 (13)	0.4489 (2)	0.48773 (7)	0.0294 (3)
H11A	0.6463	0.5716	0.4923	0.044*
H11B	0.7750	0.4449	0.5321	0.044*
H11C	0.6594	0.3077	0.4833	0.044*
C12	0.88222 (13)	0.3315 (2)	0.41201 (8)	0.0316 (3)
H12A	0.9080	0.3512	0.3621	0.047*
H12B	0.8538	0.1774	0.4179	0.047*
H12C	0.9581	0.3648	0.4498	0.047*

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\hat{A}^2)

Atomic displacement parameters $(Å^2)$

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0232 (6)	0.0284 (7)	0.0245 (6)	-0.0007 (5)	0.0069 (5)	-0.0014 (5)
N1	0.0243 (5)	0.0296 (6)	0.0250 (5)	-0.0045 (4)	0.0048 (4)	-0.0056 (4)
N2	0.0231 (5)	0.0277 (6)	0.0260 (5)	0.0025 (4)	0.0068 (4)	0.0028 (4)
01	0.0264 (5)	0.0303 (6)	0.0454 (6)	-0.0025 (4)	0.0043 (4)	-0.0106 (4)
O2	0.0247 (5)	0.0379 (6)	0.0275 (5)	0.0043 (4)	0.0092 (4)	0.0068 (4)
C2	0.0233 (6)	0.0271 (7)	0.0190 (5)	-0.0009 (5)	0.0048 (4)	0.0014 (5)
C3	0.0213 (6)	0.0344 (7)	0.0242 (6)	-0.0032 (5)	0.0019 (5)	0.0042 (5)
C4	0.0210 (6)	0.0391 (8)	0.0332 (7)	0.0037 (6)	0.0048 (5)	0.0050 (6)
C5	0.0270 (7)	0.0311 (8)	0.0377 (7)	0.0064 (6)	0.0073 (5)	-0.0012 (6)
C6	0.0245 (6)	0.0261 (7)	0.0289 (6)	0.0002 (5)	0.0048 (5)	-0.0009 (5)

supplementary materials

C7	0.0206 (6)	0.0258 (6)	0.0193 (5)	-0.0001 (5)	0.0044 (4)	0.0033 (5)	
C8	0.0200 (6)	0.0251 (6)	0.0211 (5)	0.0004 (5)	0.0058 (4)	0.0009 (5)	
C9	0.0233 (6)	0.0244 (7)	0.0219 (5)	-0.0004 (5)	0.0059 (4)	0.0006 (5)	
C10	0.0180 (5)	0.0249 (7)	0.0243 (6)	-0.0019 (5)	0.0031 (4)	-0.0005 (5)	
C11	0.0345 (7)	0.0311 (7)	0.0241 (6)	0.0005 (6)	0.0088 (5)	0.0016 (5)	
C12	0.0255 (7)	0.0323 (8)	0.0378 (7)	0.0062 (5)	0.0074 (5)	0.0059 (6)	

Geometric parameters (Å, °)

C1—N1	1.3471 (16)	C4—H4	0.9500
C1—C8	1.3858 (18)	C5—C6	1.3842 (17)
C1—H1	0.9500	С5—Н5	0.9500
N1—C2	1.3874 (16)	C6—C7	1.3936 (18)
N1—H1A	0.8800	С6—Н6	0.9500
N2-C10	1.3287 (16)	C7—C8	1.4457 (16)
N2-C12	1.4591 (16)	C8—C9	1.4380 (17)
N2-C11	1.4602 (15)	C9—C10	1.5298 (17)
O1—C9	1.2244 (16)	C11—H11A	0.9800
O2-C10	1.2431 (15)	C11—H11B	0.9800
C2—C3	1.3906 (17)	C11—H11C	0.9800
C2—C7	1.4110 (17)	C12—H12A	0.9800
C3—C4	1.380 (2)	C12—H12B	0.9800
С3—Н3	0.9500	C12—H12C	0.9800
C4—C5	1.399 (2)		
N1—C1—C8	110.00 (11)	C6—C7—C2	119.39 (11)
N1-C1-H1	125.0	C6—C7—C8	134.46 (12)
C8—C1—H1	125.0	C2—C7—C8	106.09 (11)
C1—N1—C2	109.43 (10)	C1—C8—C9	125.50 (11)
C1—N1—H1A	125.3	C1—C8—C7	106.58 (10)
C2—N1—H1A	125.3	C9—C8—C7	127.92 (12)
C10—N2—C12	120.07 (10)	O1—C9—C8	124.97 (12)
C10—N2—C11	123.31 (11)	O1—C9—C10	118.72 (11)
C12—N2—C11	116.46 (10)	C8—C9—C10	116.23 (11)
N1—C2—C3	129.53 (12)	O2-C10-N2	124.03 (11)
N1-C2-C7	107.88 (10)	O2—C10—C9	118.05 (11)
C3—C2—C7	122.53 (12)	N2-C10-C9	117.91 (10)
C4—C3—C2	116.96 (12)	N2-C11-H11A	109.5
С4—С3—Н3	121.5	N2-C11-H11B	109.5
С2—С3—Н3	121.5	H11A—C11—H11B	109.5
C3—C4—C5	121.36 (12)	N2-C11-H11C	109.5
C3—C4—H4	119.3	H11A—C11—H11C	109.5
С5—С4—Н4	119.3	H11B—C11—H11C	109.5
C6—C5—C4	121.63 (13)	N2-C12-H12A	109.5
С6—С5—Н5	119.2	N2-C12-H12B	109.5
C4—C5—H5	119.2	H12A—C12—H12B	109.5
C5—C6—C7	118.12 (12)	N2-C12-H12C	109.5
С5—С6—Н6	120.9	H12A—C12—H12C	109.5
С7—С6—Н6	120.9	H12B—C12—H12C	109.5

C8-C1-N1-C2	-1.24(14)	C6-C7-C8-C1	177 34 (13)
$C_1 $ N1 $C_2 $ C_2	175 80 (12)	C_{1} C_{2} C_{3} C_{4} C_{6} C_{1}	0.22(12)
CI = NI = C2 = C3	-175.89 (12)	C2-C/-C8-C1	0.23 (13)
C1—N1—C2—C7	1.37 (13)	C6—C7—C8—C9	-2.1 (2)
N1-C2-C3-C4	177.12 (12)	C2—C7—C8—C9	-179.18 (11)
C7—C2—C3—C4	0.20 (18)	C1C8C9O1	175.49 (12)
C2—C3—C4—C5	0.64 (19)	C7—C8—C9—O1	-5.2 (2)
C3—C4—C5—C6	-0.6 (2)	C1C8C10	-1.37 (17)
C4—C5—C6—C7	-0.3 (2)	C7—C8—C9—C10	177.94 (11)
C5—C6—C7—C2	1.11 (18)	C12—N2—C10—O2	-2.49 (19)
C5—C6—C7—C8	-175.71 (13)	C11—N2—C10—O2	-177.78 (12)
N1-C2-C7-C6	-178.59 (11)	C12—N2—C10—C9	178.11 (11)
C3—C2—C7—C6	-1.10 (18)	C11—N2—C10—C9	2.82 (18)
N1-C2-C7-C8	-0.96 (13)	O1—C9—C10—O2	-88.55 (15)
C3—C2—C7—C8	176.54 (11)	C8—C9—C10—O2	88.51 (14)
N1—C1—C8—C9	-179.96 (11)	O1-C9-C10-N2	90.89 (15)
N1—C1—C8—C7	0.61 (13)	C8—C9—C10—N2	-92.05 (14)

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	$D^{\dots}A$	<i>D</i> —H··· <i>A</i>
N1—H1A···O2 ⁱ	0.88	1.92	2.7885 (14)	170

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