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4-Diethylamino-3,5-diisopropylbenzaldehyde

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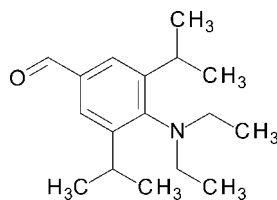
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Key indicators: single-crystal X-ray study; $T = 173$ K; mean $\sigma(\text{C}-\text{C}) = 0.003$ Å; disorder in main residue; R factor = 0.057; wR factor = 0.186; data-to-parameter ratio = 13.1.

The title benzaldehyde, $\text{C}_{17}\text{H}_{27}\text{NO}$, was prepared *via* lithiation of bromoaniline and reaction with DMF. In the crystal, the molecule adopts a C_2 -symmetrical conformation; nevertheless, two modes of disorder are present: the orientation of the aldehyde group (occupancy ratio 0.5:0.5) and of symmetry-equivalent ethyl groups [occupancy ratio 0.595 (7):0.405 (7)]. The phenylene ring and the carbonyl group are essentially coplanar [$\text{C}-\text{C}-\text{O}$ torsion angle = -179.0 (4) $^\circ$] but the dihedral angle between the mean planes of the phenylene ring and the amino group = 67.5 (2) $^\circ$. This and the long [1.414 (3) Å] aniline $\text{C}-\text{N}$ bond indicate electronic decoupling between the carbonyl and amino groups. The angle sum of 359.9 (2) $^\circ$ around the N atom results from steric compression-induced rehybridization.

Related literature

The title compound was prepared as an intermediate in the synthesis of highly solvatochromic (Detert *et al.*, 2002; Detert & Schmitt, 2006) or acidochromic fluorophores (Schmitt *et al.*, 2008, 2011). For crystal structures of anilines with a *p*-acceptor substituent, see: Fischer *et al.* (2011); Moschel *et al.* (2011). Acceptor-substituted anilines display dual fluorescence due to the formation of TICT (twisted intramolecular charge-transfer) states, see: Rotkiewicz *et al.* (1973); Okada *et al.* (1999). For the crystal structure of 4-dimethylamino-benzaldehyde, see: Gao & Zhu (2008).



Experimental

Crystal data

$\text{C}_{17}\text{H}_{27}\text{NO}$
 $M_r = 261.40$
 Monoclinic, C_2/c
 $a = 11.8061$ (9) Å
 $b = 14.3419$ (7) Å
 $c = 10.7891$ (8) Å
 $\beta = 118.478$ (3) $^\circ$

$V = 1605.78$ (19) Å³
 $Z = 4$
 Cu $K\alpha$ radiation
 $\mu = 0.50$ mm⁻¹
 $T = 173$ K
 $0.50 \times 0.20 \times 0.05$ mm

Data collection

Enraf-Nonius CAD-4
 diffractometer
 1607 measured reflections
 1533 independent reflections

1165 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.065$
 60 standard reflections every 60 min
 intensity decay: 3%

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.057$
 $wR(F^2) = 0.186$
 $S = 1.07$
 1533 reflections

117 parameters
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.31$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.23$ e Å⁻³

Data collection: *CAD-4 Software* (Enraf-Nonius, 1989); cell refinement: *CAD-4 Software*; data reduction: *CORINC* (Dräger & Gattow, 1971); program(s) used to solve structure: *SIR97* (Altomare *et al.*, 1999); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *PLATON* (Spek, 2009); software used to prepare material for publication: *PLATON*.

The authors are grateful to Heinz Kolshorn for the NMR spectra and invaluable discussions.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: BT5711).

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supporting information

Acta Cryst. (2011). E67, o3336 [https://doi.org/10.1107/S1600536811047672]

4-Diethylamino-3,5-diisopropylbenzaldehyde

Christoph Wink, Dieter Schollmeyer and Heiner Detert

S1. Comment

The aldehyde forms needle-shaped crystals of the monoclinic space group $C2/c$. The compound crystallizes in parallel layers composed of geometrical dimers with $C2$ symmetry and $z = 2$. Two modes of disorder are present. A symmetry axis through C11—C4—C1—N4 results in a $C2$ -symmetry of the molecule - but the carboxaldehyde group is disordered (50/50). The symmetry equivalent ethyl groups are also disordered but the s.o.f. is 60/40. The planes of the amino group and the benzene ring open an angle of $67.3(3)^\circ$ whereas the carbonyl group and the aromatic system are essentially coplanar (torsion angle C3—C4—C11—O12: $-179.0(4)^\circ$). The sum of the bond angles around the nitrogen atom in both conformers is $359.9(2)^\circ$, corresponding to a sp^2 hybridization. With $1.414(3) \text{ \AA}$, the aniline C—N bond is much longer than the corresponding bond ($1.366(2) \text{ \AA}$) in the sterically undisturbed dimethylaminobenzaldehyde (Gao & Zhu, 2008). This may be in part due to steric congestion, but also due to inhibited conjugation. Correspondingly, the carbonyl bond length of $1.114(4) \text{ \AA}$ in the title compound is shorter than in the dimethylamino derivative ($1.204(2) \text{ \AA}$, $1.212(3) \text{ \AA}$) and with $1.470(4) \text{ \AA}$ the aryl-carbonyl bond C4—C11 is longer than in the reference compound ($1.457(3) \text{ \AA}$, $1.454(2) \text{ \AA}$).

S2. Experimental

4-Bromo-*N,N*-diethyl-2,6-diisopropylaniline (0.76 g, 2.5 mmol) was dissolved in THF (15 ml) under nitrogen in a flame-dried Schlenk tube. The solution was cooled to 195 K and *n*-BuLi (2.5 M in heptane, 1 ml) was added dropwise. After stirring for 1 h, dry DMF (0.2 ml, 2.5 mmol) was added carefully, stirring was continued for 15 min at 195 K, and the solution was allowed to reach room temperature. Aqueous NH_4Cl (conc. 15 ml) was added and the mixture extracted with ethyl acetate (3 * 20 ml). The pooled organic solutions were dried (Na_2SO_4), concentrated *in vacuo* and the product was isolated *via* column chromatography (SiO_2 , petroleum ether / ethyl acetate = 15 / 1) $R_f = 0.4$ (petroleum ether / ethyl acetate = 9 / 1). The aldehyde was isolated as a yellowish oil that crystallized upon standing for several days. Yield: 0.23 g (35%)

S3. Refinement

Hydrogen atoms attached to carbons were placed at calculated positions with C—H = 0.95 \AA (aromatic) or $0.98\text{--}0.99 \text{ \AA}$ (sp^3 C-atom). All H atoms were refined in the riding-model approximation with isotropic displacement parameters set at 1.2–1.5 times of the U_{eq} of the parent atom.

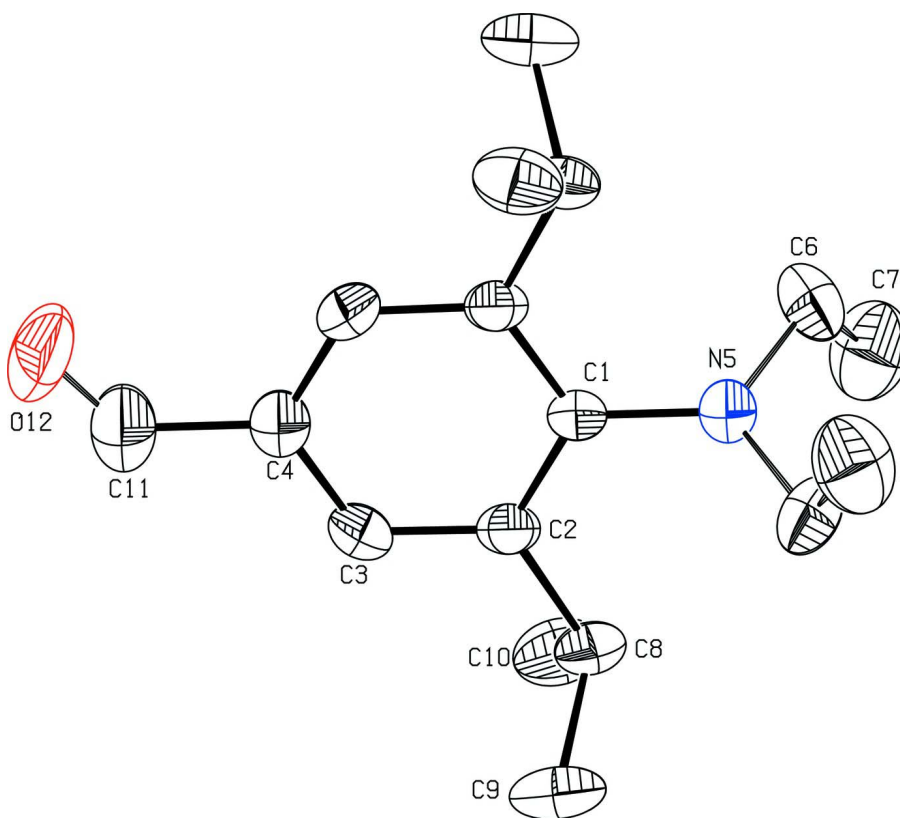


Figure 1

View of compound I. Displacement ellipsoids are drawn at the 50% probability level. Only major conformer is shown.

4-Diethylamino-3,5-diisopropylbenzaldehyde

Crystal data

$C_{17}H_{27}NO$
 $M_r = 261.40$
 Monoclinic, $C2/c$
 Hall symbol: $-C\ 2yc$
 $a = 11.8061\ (9)\ \text{\AA}$
 $b = 14.3419\ (7)\ \text{\AA}$
 $c = 10.7891\ (8)\ \text{\AA}$
 $\beta = 118.478\ (3)^\circ$
 $V = 1605.78\ (19)\ \text{\AA}^3$
 $Z = 4$

$F(000) = 576$
 $D_x = 1.081\ \text{Mg m}^{-3}$
 Cu $K\alpha$ radiation, $\lambda = 1.54178\ \text{\AA}$
 Cell parameters from 25 reflections
 $\theta = 60\text{--}70^\circ$
 $\mu = 0.50\ \text{mm}^{-1}$
 $T = 173\ \text{K}$
 Needle, colourless
 $0.50 \times 0.20 \times 0.05\ \text{mm}$

Data collection

Enraf–Nonius CAD-4
 diffractometer
 Radiation source: rotating anode
 Graphite monochromator
 $\omega/2\theta$ scans
 1607 measured reflections
 1533 independent reflections
 1165 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.065$
 $\theta_{\text{max}} = 70.0^\circ$, $\theta_{\text{min}} = 5.3^\circ$
 $h = 0 \rightarrow 14$
 $k = 0 \rightarrow 17$
 $l = -13 \rightarrow 11$
 60 standard reflections every 60 min
 intensity decay: 3%

Refinement

Refinement on F^2

Least-squares matrix: full

 $R[F^2 > 2\sigma(F^2)] = 0.057$ $wR(F^2) = 0.186$ $S = 1.07$

1533 reflections

117 parameters

0 restraints

Primary atom site location: structure-invariant
direct methodsSecondary atom site location: difference Fourier
mapHydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

 $w = 1/[\sigma^2(F_o^2) + (0.1003P)^2 + 0.8224P]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{\max} < 0.001$ $\Delta\rho_{\max} = 0.31 \text{ e } \text{\AA}^{-3}$ $\Delta\rho_{\min} = -0.23 \text{ e } \text{\AA}^{-3}$ Extinction correction: *SHELXL97* (Sheldrick,
2008), $F_c^* = kF_c[1 + 0.001xF_c^2\lambda^3/\sin(2\theta)]^{-1/4}$

Extinction coefficient: 0.0061 (10)

Special details

Experimental. $^1\text{H-NMR}$ (400 MHz, CDCl_3): 9.95 (s, 1 H, CHO), 6.61 (s, 2 H, 2-H, 6-H), 3.49 (sept, $^3J = 6.9$ Hz, 2 H, CH (i-Pr)), 3.11 (q, $^3J = 7.1$ Hz, 4 H, N- CH_2), 1.22 (d, $^3J = 6.9$ Hz, 12 H, CH_3 (iPr)), 1.04 (t, $^3J = 7.1$ Hz, 6 H, CH_3 (Et)). $^{13}\text{C-NMR}$ (75 MHz, CDCl_3): 192.5 (CHO), 152.3 (C-4), 150.9 (C-3, C-5), 134.3 (C-1), 126.0 (C-2, C-6), 48.9 (N- CH_2), 29.2 (CH (iPr)), 24.5 (CH_3 (iPr)), 15.2 (CH_3 (Et)).IR (ATR) $\nu = 2963, 2928, 2869, 2723, 1696, 1594, 1568, 1457, 1365, 1268, 1170, 1104, 1065, 941, 892, 783, 724 \text{ cm}^{-1}$.HR-ESI-MS: found 262.2177, calc. 262.2171 for $(\text{M}+\text{H}^+)$.**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	$U_{\text{iso}}^*/U_{\text{eq}}$	Occ. (<1)
C1	0.5000	0.29898 (16)	0.2500	0.0301 (6)	
C2	0.49043 (16)	0.25001 (12)	0.13197 (17)	0.0329 (5)	
C3	0.49237 (18)	0.15335 (13)	0.13533 (18)	0.0407 (5)	
H3	0.4884	0.1198	0.0574	0.049*	
C4	0.5000	0.10460 (18)	0.2500	0.0445 (7)	
N5	0.5000	0.39757 (14)	0.2500	0.0394 (6)	
C6	0.3927 (4)	0.4517 (2)	0.2464 (5)	0.0477 (12)	0.595 (7)
H6A	0.4284	0.5063	0.3095	0.057*	0.595 (7)
H6B	0.3471	0.4125	0.2841	0.057*	0.595 (7)
C7	0.2975 (5)	0.4852 (3)	0.1028 (5)	0.0704 (16)	0.595 (7)
H7A	0.3405	0.5274	0.0668	0.106*	0.595 (7)
H7B	0.2275	0.5186	0.1082	0.106*	0.595 (7)
H7C	0.2620	0.4318	0.0391	0.106*	0.595 (7)
C6A	0.3813 (6)	0.4420 (4)	0.1519 (7)	0.0511 (19)	0.405 (7)
H6C	0.3242	0.3953	0.0827	0.061*	0.405 (7)
H6D	0.3999	0.4910	0.0997	0.061*	0.405 (7)
C7A	0.3127 (8)	0.4849 (6)	0.2250 (9)	0.084 (3)	0.405 (7)
H7D	0.2754	0.4355	0.2570	0.126*	0.405 (7)

H7E	0.2439	0.5260	0.1593	0.126*	0.405 (7)
H7F	0.3741	0.5212	0.3064	0.126*	0.405 (7)
C8	0.47720 (17)	0.29910 (13)	0.00066 (18)	0.0384 (5)	
H8	0.4765	0.3678	0.0159	0.046*	
C9	0.5913 (2)	0.27726 (19)	-0.0249 (2)	0.0577 (7)	
H9A	0.6715	0.2966	0.0573	0.087*	
H9B	0.5812	0.3111	-0.1085	0.087*	
H9C	0.5941	0.2101	-0.0399	0.087*	
C10	0.3504 (2)	0.27324 (18)	-0.1286 (2)	0.0535 (6)	
H10A	0.3516	0.2070	-0.1502	0.080*	
H10B	0.3397	0.3108	-0.2094	0.080*	
H10C	0.2786	0.2853	-0.1093	0.080*	
C11	0.5000	0.0021 (2)	0.2500	0.0730 (11)	
H11	0.4960	-0.0251	0.1677	0.088*	0.50
O12	0.5039 (6)	-0.0493 (2)	0.3294 (5)	0.1012 (16)	0.50

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0306 (11)	0.0327 (12)	0.0323 (12)	0.000	0.0193 (9)	0.000
C2	0.0347 (9)	0.0379 (10)	0.0326 (9)	0.0009 (7)	0.0213 (7)	0.0008 (6)
C3	0.0542 (11)	0.0377 (10)	0.0404 (10)	0.0018 (8)	0.0309 (9)	-0.0059 (7)
C4	0.0589 (17)	0.0334 (14)	0.0487 (15)	0.000	0.0317 (13)	0.000
N5	0.0500 (13)	0.0301 (11)	0.0438 (12)	0.000	0.0272 (10)	0.000
C6	0.051 (2)	0.0404 (19)	0.046 (2)	0.0115 (16)	0.0187 (17)	-0.0043 (15)
C7	0.062 (3)	0.062 (3)	0.073 (3)	0.008 (2)	0.020 (2)	0.017 (2)
C6A	0.063 (4)	0.037 (3)	0.054 (4)	0.012 (3)	0.028 (3)	0.008 (2)
C7A	0.068 (5)	0.082 (5)	0.092 (6)	0.020 (4)	0.031 (4)	-0.032 (4)
C8	0.0425 (10)	0.0476 (11)	0.0311 (9)	-0.0020 (8)	0.0225 (8)	0.0020 (7)
C9	0.0548 (13)	0.0874 (17)	0.0472 (12)	-0.0001 (11)	0.0374 (10)	0.0050 (11)
C10	0.0519 (12)	0.0748 (15)	0.0329 (10)	-0.0049 (10)	0.0194 (9)	0.0035 (9)
C11	0.112 (3)	0.0387 (17)	0.083 (3)	0.000	0.059 (2)	0.000
O12	0.195 (5)	0.0416 (19)	0.101 (3)	-0.003 (2)	0.098 (3)	0.0176 (19)

Geometric parameters (Å, °)

C1—C2	1.411 (2)	C6A—C7A	1.505 (10)
C1—C2 ⁱ	1.411 (2)	C6A—H6C	0.9900
C1—N5	1.414 (3)	C6A—H6D	0.9900
C2—C3	1.387 (3)	C7A—H7D	0.9800
C2—C8	1.522 (2)	C7A—H7E	0.9800
C3—C4	1.386 (2)	C7A—H7F	0.9800
C3—H3	0.9500	C8—C10	1.527 (3)
C4—C3 ⁱ	1.386 (2)	C8—C9	1.530 (3)
C4—C11	1.470 (4)	C8—H8	1.0000
N5—C6A ⁱ	1.441 (6)	C9—H9A	0.9800
N5—C6A	1.441 (6)	C9—H9B	0.9800
N5—C6 ⁱ	1.470 (4)	C9—H9C	0.9800

N5—C6	1.470 (4)	C10—H10A	0.9800
C6—C7	1.495 (6)	C10—H10B	0.9800
C6—H6A	0.9900	C10—H10C	0.9800
C6—H6B	0.9900	C11—O12 ⁱ	1.114 (4)
C7—H7A	0.9800	C11—O12	1.114 (4)
C7—H7B	0.9800	C11—H11	0.9500
C7—H7C	0.9800		
C2—C1—C2 ⁱ	120.3 (2)	C7A—C6A—H6D	109.2
C2—C1—N5	119.86 (11)	H6C—C6A—H6D	107.9
C2 ⁱ —C1—N5	119.86 (11)	C6A—C7A—H7D	109.5
C3—C2—C1	118.74 (16)	C6A—C7A—H7E	109.5
C3—C2—C8	118.68 (15)	H7D—C7A—H7E	109.5
C1—C2—C8	122.58 (17)	C6A—C7A—H7F	109.5
C4—C3—C2	121.39 (17)	H7D—C7A—H7F	109.5
C4—C3—H3	119.3	H7E—C7A—H7F	109.5
C2—C3—H3	119.3	C2—C8—C10	111.02 (15)
C3 ⁱ —C4—C3	119.4 (2)	C2—C8—C9	111.38 (15)
C3 ⁱ —C4—C11	120.29 (12)	C10—C8—C9	110.43 (16)
C3—C4—C11	120.29 (12)	C2—C8—H8	108.0
C1—N5—C6A ⁱ	116.2 (2)	C10—C8—H8	108.0
C1—N5—C6A	116.2 (2)	C9—C8—H8	108.0
C6A ⁱ —N5—C6A	127.5 (5)	C8—C9—H9A	109.5
C1—N5—C6 ⁱ	121.86 (16)	C8—C9—H9B	109.5
C6A—N5—C6 ⁱ	108.1 (3)	H9A—C9—H9B	109.5
C1—N5—C6	121.86 (16)	C8—C9—H9C	109.5
C6A ⁱ —N5—C6	108.1 (3)	H9A—C9—H9C	109.5
C6 ⁱ —N5—C6	116.3 (3)	H9B—C9—H9C	109.5
N5—C6—C7	114.2 (4)	C8—C10—H10A	109.5
N5—C6—H6A	108.7	C8—C10—H10B	109.5
C7—C6—H6A	108.7	H10A—C10—H10B	109.5
N5—C6—H6B	108.7	C8—C10—H10C	109.5
C7—C6—H6B	108.7	H10A—C10—H10C	109.5
H6A—C6—H6B	107.6	H10B—C10—H10C	109.5
N5—C6A—C7A	112.0 (6)	O12 ⁱ —C11—C4	131.4 (3)
N5—C6A—H6C	109.2	O12—C11—C4	131.4 (3)
C7A—C6A—H6C	109.2	O12—C11—H11	114.3
N5—C6A—H6D	109.2	C4—C11—H11	114.3
C2 ⁱ —C1—C2—C3	−0.85 (12)	C1—N5—C6—C7	−97.8 (3)
N5—C1—C2—C3	179.15 (12)	C6A ⁱ —N5—C6—C7	123.5 (4)
C2 ⁱ —C1—C2—C8	178.76 (17)	C6A—N5—C6—C7	−4.4 (4)
N5—C1—C2—C8	−1.24 (17)	C6 ⁱ —N5—C6—C7	82.2 (3)
C1—C2—C3—C4	1.7 (2)	C1—N5—C6A—C7A	107.0 (5)
C8—C2—C3—C4	−177.89 (13)	C6A ⁱ —N5—C6A—C7A	−73.0 (5)
C2—C3—C4—C3 ⁱ	−0.89 (12)	C6 ⁱ —N5—C6A—C7A	−111.7 (5)
C2—C3—C4—C11	179.11 (12)	C6—N5—C6A—C7A	−2.0 (4)
C2—C1—N5—C6A ⁱ	−112.2 (3)	C3—C2—C8—C10	61.2 (2)

C2 ⁱ —C1—N5—C6A ⁱ	67.8 (3)	C1—C2—C8—C10	-118.38 (18)
C2—C1—N5—C6A	67.8 (3)	C3—C2—C8—C9	-62.3 (2)
C2 ⁱ —C1—N5—C6A	-112.2 (3)	C1—C2—C8—C9	118.10 (18)
C2—C1—N5—C6 ⁱ	-67.7 (2)	C3 ⁱ —C4—C11—O12 ⁱ	-179.0 (4)
C2 ⁱ —C1—N5—C6 ⁱ	112.3 (2)	C3—C4—C11—O12 ⁱ	1.0 (4)
C2—C1—N5—C6	112.3 (2)	C3 ⁱ —C4—C11—O12	1.0 (4)
C2 ⁱ —C1—N5—C6	-67.7 (2)	C3—C4—C11—O12	-179.0 (4)

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